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**EFFECTS OF CIGARETTE SMOKE
ON KILLER CELL ACTIVATION IN
CHRONIC OBSTRUCTIVE PULMONARY DISEASE**

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Dedication

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Abbreviations

AAT	α_1 -antitrypsin
ADCC	antibody-dependent cellular cytotoxicity
APCs	antigen-presenting cells
ATS	American thoracic society
BAL	bronchoalveolar lavage
BLyS	TNF-related ligand B-lymphocyte stimulator
BODE	body-mass index (B), the degree of airflow obstruction (O), dyspnoea (D), and exercise capacity (E)
BSA	bovine serum albumin
CCR	C-C chemokine receptor
COPD	chronic obstructive pulmonary disease
CSE	cigarette smoke extract
cuS-COPD	current smokers with COPD
DCs	dendritic cells
DTT	2,3-dihydroxybutane-1,4-dithiol (dithiotheritol)
ECD	phycoerythrin-Texas Red-x
ECP	eosinophil cationic protein
EDTA	ethylenediaminetetraacetic acid
EGF	epidermal growth factor

Egr-1	early growth response gene 1
ENA-78	epithelial neutrophils activating protein of 78 kDa
EPO	eosinophil peroxidase
ET-1	endothelin-1
exS-COPD	ex-smokers with COPD
FACS	fluorescent activated cell sorting
Fas-L	Fas ligand
FCS	Foetal calf serum
FEV ₁	forced expiratory volume in one second
FGF	fibroblast growth factors
FITC	fluorescein isothiocyanate
FMO	fluorescence minus one
FVC	forced vital capacity
α -Galcer	α -galactosylceramide
G-CSF	granulocyte colony-stimulating factor
GM-CSF	granulocyte-macrophage colony stimulating factor
GOLD	global initiative for chronic obstructive lung disease
GPCRs	G-protein-coupled receptors
GRO- α	growth-related oncogene- α
4-HNE	4-hydroxy-2-nonenal-modified
HNS	healthy non-smokers

HS	current healthy smokers
ICAM-1	intercellular adhesion molecule 1
iNKT	invariant natural killer T cell
iNOS	inducible NO synthase
IP-10	interferon-inducible protein-10
ITAM	immunoreceptor tyrosine-based activation motif
ITIM	immunotyrosine-based inhibitory motif
KIRs	killer cell immunoglobulin-like receptors
LCs	Langerhans cells
LFA-1	lymphocyte function-associated antigen 1
LILRs	leukocyte immunoglobulin (Ig)-like receptors
LTB ₄	leukotriene B ₄
LPS	lipopolysaccharides
MAC	macrophage antigen
MAPK	mitogen-activated protein kinase
MC _T	mucosal mast cells
MC _{TC}	connective tissue mast cells
mDCs	myeloid dendritic cells
MHC	major histocompatibility complex
MIG	monokine induced by interferon-gamma
MMP	matrix metalloproteinase

MMPS	matrix metalloprotease enzymes
MPO	myeloperoxidase
MRC	Medical Research Council
NE	neutrophil elastase
NF- κ B	nuclear factor- κ B
NK	natural killer
NKT	natural killer T cell
NO	nitric oxide
PBMCs	peripheral blood mononuclear cell
PBS	phosphate buffered saline
PC5	phycoerythrin-cyanin 5.1
PC7	phycoerythrin-cyanin 7
pDCs	plasmacytoid dendritic cells
PE	phycoerythrin
PGE ₂	prostaglandin E ₂
PGF _{2α}	prostaglandin F _{2α}
RANTES	released by activated normal T cells expressed and secreted
REC	Research Ethics Committee
ROS	reactive oxygen species
RPMI 1640	Roswell Park Memorial Institute media

RV	rhinovirus
TARC	thymus- and activation-dependent chemokine
TB	tuberculosis
Tc	cytotoxic T cell
TCR	T cell receptor
TEER	trans-epithelial electric resistance
TGF- β	transforming growth factor- β
Th	helper T cell
TIMPs	tissue inhibitors of MMPs
TLA-1	T-cell intra-cytoplasmic antigen-1
TNF	tumour-necrosis factor
Treg	regulatory T cells
TSLP	thymic stromal lymphopoietin
VEGF	vascular endothelial growth factor
WHO	World Health Organization

Publications Arising from this Work

1. **J Wang**, Richard A Urbanowicz, Patrick J. Tighe, Ian Todd, Jonathan M. Corne, and Lucy C. Fairclough. Differential Activation of Killer Cells in the Circulation and the Lung: A Study of Current Smoking Status and Chronic Obstructive Pulmonary Disease (COPD). PLoS ONE. 2013 Mar; 8(3): e58556.
2. **J Wang**, Ian Todd, Jonathan M. Corne, Patrick J. Tighe, and Lucy C. Fairclough. Cigarette smoke extract suppresses NK cells activation in current smokers with COPD. Eur Respir J. 2014; in preparation

Poster Presentations Arising from this Work

1. **J Wang**, Richard A Urbanowicz, Jonathan M. Corne, Ian Todd, and Lucy C. Fairclough. Cytotoxic Cells in Bronchoalveolar Lavage (BAL) Fluid from COPD patients. ATS conference 2009; San Diego, USA.

Abstract

Chronic Obstructive Pulmonary Disease (COPD) is a chronic inflammatory disease involving both innate and adaptive immune responses. Abnormal numbers of inflammatory cells have been examined in COPD subjects, as well as the effects of cigarette smoking on immune cells and molecules. Killer cells, including CD8⁺ T cells, NKT-like cells and NK cells, are thought to play a role in the development of COPD through their cytotoxic functions. In this project, we report *ex vivo*, activation levels of these cell types in COPD patients, as well as effects induced by cigarette smoke extract *in vitro*.

PBMCs were collected from healthy non-smokers (HNS), current healthy smokers (HS), current smokers with COPD (cuS-COPD) and ex-smokers with COPD (exS-COPD). Activation levels of interest and CSE effects on them were analysed by flow cytometry.

Killer cells, including CD8⁺ T cells, NKT-like cells and NK cells, were significantly activated in current smokers with or without COPD compared to healthy non-smokers. Furthermore, KIR (CD158e1) expression was dramatically lower in smokers with or without COPD in comparison with healthy non-smokers. The cytotoxicity of CD8⁺ T cells from both current smokers and ex-smokers with COPD patients were significantly less than that in healthy volunteers. Also, *in vitro*, CSE markedly decreased

IL-15 treated NK cell activation in current smokers with COPD compared to other three groups. The expression of granzyme B was also significantly inhibited on IL-15 stimulated NK cells when CSE was added.

We conclude systemic *ex vivo* killer cell activation is smoking rather than disease related. Cigarette smoking has immunosuppressive effects on killer cell activation and granzyme B expression in PBMCs from current smokers with COPD.

Introduction

1.1 Chronic Obstructive Pulmonary Disease

1.1.1 Definition, Prevalence and Global burden

COPD, as a common, preventable and treatable disease, is characterised as a chronic inflammatory disease causing airway limitation that is usually progressive but not completely reversible (Celli and MacNee, 2004). It is the fourth leading worldwide cause of morbidity and mortality, and is predicted to become the third most frequent burden of disease in the world by 2030, after heart disease and stroke (Rabe et al., 2007). The estimated death rate of COPD shows geographical differences. In the US, according to updated statistics from the Centre for Disease Control (CDC), 0.04% of people died from COPD in 2009. Compared with the decreasing mortality in some diseases, such as heart disease and stroke, COPD is the only major disease with an increasing death rate. In European countries, mortality from COPD varies from 0.025% to 0.075% of the adult population. In Japan, the reported mortality from COPD is 0.04%, whereas nearly 1.3% of deaths result from COPD in China (Buist et al., 2007, Fukuchi et al., 2004, Halbert et al., 2006). Gender differences are also seen: morbidity from COPD is higher in men than in women in the UK (Soriano et al., 2000). Nevertheless, some studies indicate that the prevalence of COPD seems almost equal in men and women and that women are more susceptible to

COPD than men (Xu et al., 1994, Silverman et al., 2000). Medical costs of COPD are an economic burden in both developed countries and developing countries (Chapman et al., 2006). There is a positive correlation between the severity of COPD and the cost of care (Jansson et al., 2002), although costs per patient for COPD are variable across countries due to how health care is provided and paid .

1.1.2 Risk factors

Prevalence estimates of the disorder show considerable variability across populations, suggesting that risk factors can affect populations differently. Although only 15% to 20% of smokers develop clinical COPD (Brown et al., 1994), cigarette smoking is still the common and key pathological driver in the development and progression of COPD (Burrows et al., 1977). The Fletcher curve (Figure 1.1) shows the importance of smoking cessation in patients with COPD.

Besides, other factors are involved. Genetic predisposition now also seems to play an important role in the development of COPD. α_1 -antitrypsin (AAT), a major circulating inhibitor of serine proteases, is the best established antiprotease associated with COPD (Hill et al., 1999, Ishii et al., 2000, Sandhaus, 2009, Zielinski and Kuca, 2004). It has been

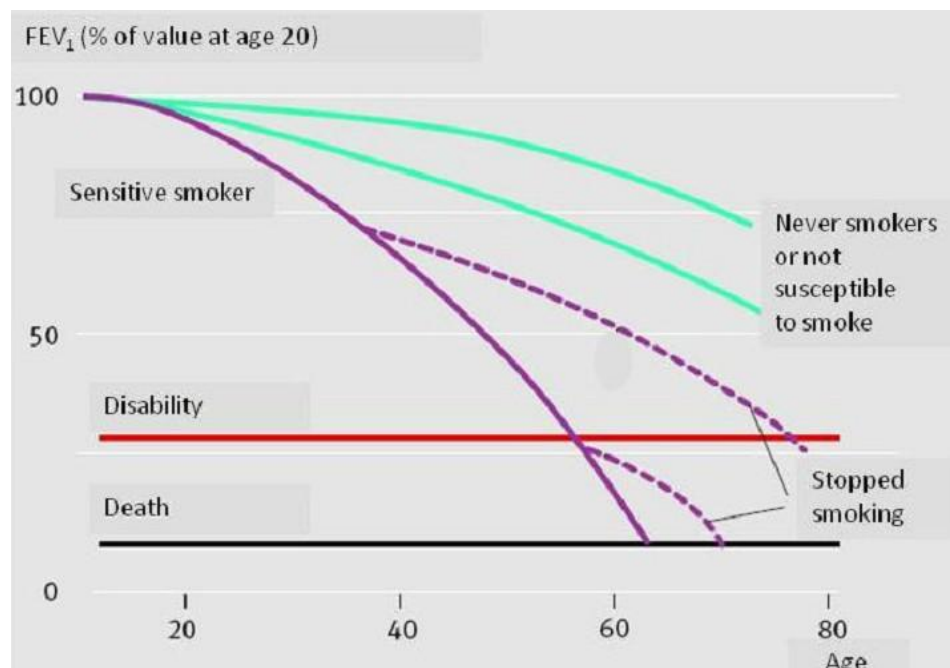


Figure 1.1 Age-related decline in FEV₁ with different smoking history. (Adapted from Fletcher et al. Definitions of emphysema, chronic bronchitis, asthma, and airflow obstruction: 25 years on from the Ciba symposium. Thorax 1984; 39: 81-85)

observed that AAT deficiency can lead to early onset emphysema in COPD patients (Ekeowa et al., 2009, Eriksson, 1965, Ranes and Stoller, 2005, Strange, 2013). Metalloproteases, proinflammatory cytokines, various antioxidant enzymes and detoxifying enzymes are also linked to COPD (Wood and Stockley, 2006). Occupational exposure to biomass smoke, environmental pollution with noxious particles and gases, chronic asthma, history of pulmonary tuberculosis and poor socioeconomic status can all result in COPD (Balmes et al., 2003, Barker et al., 1991, Ezzati, 2005,

Orozco-Levi et al., 2006, Prescott and Vestbo, 1999, Viegi and Di Pede, 2002). These may explain why there are around one-third of COPD patients who have never smoked (Salvi and Barnes, 2009).

Cigarette Smoke

Cigarette smoking has been implicated as an etiological risk factor resulting in various chronic diseases with high morbidity and mortality, including cancers, heart diseases, variety of infections and chronic lung diseases (Almirall et al., 1999, Arcavi and Benowitz, 2004, Soporì, 2002).

The smoke from a cigarette contains thousands of chemicals which have various toxic, mutagenic and carcinogenic activities (Hoffmann and Wynder, 1986, Stedman, 1968). The concentrations of chemical ingredients also vary widely in the different cigarette brands. Studies show that various components of cigarette smoke can affect the immune response in different cell types (Lee et al., 2007b, McCue et al., 2000, Ouyang et al., 2000). Two major components of smoke that leads to many of deleterious effects are nicotine and tar (Hoffmann and Wynder, 1986, Stedman, 1968). Holt *et al* has shown that cigarette smoke containing high levels of tar and nicotine is more immunosuppressive than the smoke

from low-tar low-nicotine cigarettes (Holt et al., 1976). However, effects of cigarette smoke on the immune system are biphasic, which is mainly determined by the components and duration of exposure. For example, unlike nicotine and tar, tobacco glycoprotein and metals present in cigarette smoke are generally immunostimulatory (Brooks et al., 1990, Francus et al., 1992).

Previous studies have demonstrated that cigarette smoke causes structural and functional changes in the respiratory ciliary epithelium (Cantin et al., 2006, Mio et al., 1997, Wyatt et al., 1999), lung surfactant and immune cells (Mohan, 2002a). In COPD patients, abnormal cell numbers of different cell types have been observed, including CD8⁺ T cells, alveolar macrophages, dendritic cells (DCs), neutrophils and NK cells in bronchoalveolar lavage (BAL) fluid, epithelium and adventitia of small airways, sputum or peripheral blood samples (O'Shaughnessy et al., 1997, Saetta et al., 1998, Baraldo et al., 2004, Barnes, 2004b, Demedts et al., 2007, Urbanowicz et al., 2009). Therefore, accumulating data suggested that chronic inhalation of cigarette smoke alters a wide range of immunologic function, including both innate and adaptive immune responses.

1.1.3 Systemic effects of COPD

Studies have shown that COPD is associated with both pulmonary and systemic inflammation. Weight loss, skeletal muscle dysfunction, cardiovascular disease, cancer, nutritional abnormalities and other organ effects all contribute the high burden of COPD together with respiratory symptoms protein (Agusti, 2007, Gosker et al., 2000, Pryor and Stone, 1993, Schols et al., 1996, Skillrud et al., 1986, Van den Heuvel et al., 1999), which result in a pronounced deterioration in health status of life.

1.1.4 Comparison of COPD with asthma and tuberculosis

COPD, asthma and tuberculosis (TB) are three main public health problems related to lung worldwide. Studies have shown that these diseases have common risk factors, such as cigarette smoking, low socioeconomic status and dysregulation of host defence functions. Also, they can co-exist, which result in co-morbid conditons. Table 1.1 lists the common charasteristics and differences of among these diseases.

1.1.5 Pathology of COPD

Chronic obstructive bronchiolitis (swelling and inflammation of the lining of the airways that results in narrowing and

Table 1.1 Comparison of COPD with asthma and TB

COPD	Asthma	TB
Common Features		
<ul style="list-style-type: none"> • Pulmonary diseases • All carry a significant burden in terms of morbidity and mortality worldwide • Induce inflammatory responses 		
Differences		
Epidemiology		
Worldwide prevalence; Not infectious	High prevalence in the developed countries; Not infectious	Over 95% of TB cases in developing countries; Infectious
Patients group		
Mainly adults ≥ 40 years old	Mainly young children	Mainly young adults
Symptoms		
Chronic cough ; Dyspnea; Large amounts of mucus production; Wheezing; Chest tightness	Cough (specially at night); Wheezing; Shortness of breath; Chest tightness, pain or pressure; Fatigue	Cough (usually cough up mucus); Coughing up blood; Fatigue; Fever; Excessive sweating, especially at night; Unintentional weight loss
Burden of mortality		
The fourth leading worldwide cause of morbidity and mortality	Most cases are managed at a primary care level.	Second only to HIV/AIDS as the greatest infectious killer worldwide
Disease development		
Airway limitation is not reversible; Have features of asthma	Airway limitation is reversible; Co-exist with COPD	Not all patients develop airway limitation; COPD as a sequel to active TB
Immune cells mainly involved		
Neutrophils, macrophages, CD8 ⁺ T cells, NK cells and NKT cells; Both Th1 and Th2 cell-induced immunity	Eosinophils, CD4 ⁺ T cells; Mainly Th1 cells-mediated immunity	CD4 ⁺ T cells play a crucial role; Mainly Th1 cell-induced immunity
References		
(Sciurba, 2004)	(Braman, 2006)..	(Chakrabarti et al., 2007)

obstruction of the airways), emphysema (permanent destructive enlargement of airspaces distal to the terminal bronchioles, caused by loss of lung elastic recoil) and mucus hypersecretion (productively coughing and bringing up sputum for >3 months for at least two consecutive years) are recognized as three distinct pathological features of most COPD patients (Barnes, 2000, Kim et al., 1991). They affect all parts of the lungs including parenchyma and airways and increase the resistance of the conducting airways and lung compliance. The severity of COPD is related to the remodelling of the small airways, especially of the terminal bronchioles including increased airway smooth muscle and goblet cell hyperplasia, submucosal gland hypertrophy and epithelial metaplasia (Hogg et al., 2004).

1.1.6 Classification of COPD

Currently, two staging systems have been applied to COPD. Global Initiative for Chronic Obstructive Lung Disease (GOLD) has confirmed the classification of COPD with spirometry. According to the two values of the postbronchodilator spirometric measurement, namely, forced expiratory volume in one second (FEV₁) and forced vital capacity (FVC), COPD is classified from stage 0 to stage III (Table 1.2) (www.goldcopd.org., 2011). However, COPD patients carry

Table 1.2 GOLD classification of COPD

Stage	Spirometry	Symptoms
0: At risk	Normal spirometry	Chronic symptoms (cough, sputum production)
I: Mild COPD	FEV ₁ /FVC<70% FEV ₁ greater than or equal to 80% predicted	With or without chronic symptoms (cough, sputum production)
II: Moderate COPD	FEV ₁ /FVC<70% FEV ₁ greater than or equal to 30% to 80% predicted IIa: FEV ₁ greater than or equal to 50% to 80% predicted IIb: FEV ₁ greater than or equal to 30% to 50% predicted	With or without chronic symptoms (cough, sputum production, dyspnea)
III: Severe COPD	FEV ₁ /FVC<70% FEV ₁ <30% predicted or FEV ₁ <50% plus respiratory failure or clinical signs of right heart failure Respiratory failure: arterial partial pressure of oxygen<60mm Hg with or without arterial partial pressure of carbon dioxide greater than or equal to 50mm Hg while breathing air at sea level	
Source: <i>The GOLD Workshop level</i>		

both physical impairment and mental burden related to their disease and its symptoms, such as anxiety and depression (Gudmundsson et al., 2005, Wagena et al., 2005); the classification of COPD based on GOLD stages is not sufficient for explaining those mental symptoms (Dahlen and Janson, 2002, Hill et al., 2008, Celli et al., 2004). The BODE index (Body Mass-Index, Airflow Obstruction, Dyspnea and Exercise Capacity) is a multidimensional scoring system for COPD. It comprises an assessment of symptoms, a surrogate of the nutritional state, and exercise capacity together with the spirometric measure of airflow (FEV₁) (Celli et al., 2004). It has been shown to be superior in predicting hospitalization and the mortality risk of COPD patients and better correlates to health status (Celli et al., 2004, Ong et al., 2005, Ong et al., 2006). Table 1.3 shows the BODE index classification.

To detect COPD early, the disease should be suspected when a patient has with chronic cough and sputum production and a history of COPD risk factors.

Table 1.3 BODE index for COPD

Variable	BODE index point			
	0	1	2	3
FEV₁% predicted	≥65	50-64	36-49	≤35
Distance walked in 6 minutes (m)	≥349	250-349	150-249	≤149
MRC dyspnoea scale	0-1	2	3	4
BMI (Kg/m²)	>21		≤21	

Modified MRC Dyspnoea Scale

0. Breathless only with strenuous exercise
1. Short of breath when hurrying on the level or walking up a slight hill
2. Slower than most people of the same age on the level because of breathlessness or have to stop for breath when walking atown pace on the level
3. Stop for breath after walking about 100 meters or after a few minutes at own pace on the level
4. Too breathless to leave the house or breathless when dressing

1.2 Pathological hallmarks in COPD

In the last decade, there have been many studies performed on bronchial biopsies, lung parenchyma and peripheral blood obtained from COPD patients compared with healthy non-smokers and smokers with normal lung function. They provide new insights on the role of the different inflammatory and structural cells, mediators and signalling pathways, contributing to a better knowledge of the pathogenesis of COPD. Immunological cells, such as neutrophils, macrophages, DCs, T lymphocytes, epithelial cells, NK cells, NKT-like cells and so on, are mostly involved in the abnormal inflammatory response that occurs in COPD. These cells play different roles in producing structural changes in the airways including loss of respiratory bronchioles, alveolar ducts and alveoli, which further lead to smooth muscle hypertrophy, epithelial disruption, fibrosis, mucociliary dysfunction and lung destruction. This section summarizes how these immune cells work in COPD patients.

1.2.1 Immune cells in COPD

1.2.1.1 Neutrophils

Neutrophils are the only cell type in mammals permitted to migrate to any part of the body (Nathan, 2006). As one of the front line defensive cells of the immune system, they

contribute to both tissue repair and tissue damage (Henson et al., 2006, Weiss, 1989). Once injury happens, activated neutrophils accumulate to infected tissues and release proteases, such as neutrophil elastase, cathepsin G, matrix metalloproteinases (MMPs), and reactive oxygen intermediate including superoxide, onzone, hydroxyl radical that contribute to tissue damage (Nathan, 2006). On the other hand, neutrophils produce destructive oxidizing agents and attempts at blocking neutrophil function that have resulted in significantly diminished early tissue destruction (Brickson et al., 2003). According to this unique capacity and without the serious side effects of increasing the host's risk of infection (Nathan, 2002), targeting neutrophils can be considered as an anti-inflammatory therapeutic in many diseases. Neutrophils are also life-saving decision-markers that trigger and maintain an immune response by producing molecular signals. Generated signals can activate epithelial cells, macrophages, mast cells, endothelial cells, T cells, B cells, and also neutrophils themselves. For instance, tumour-necrosis factor (TNF) produced by neutrophils drives DC and macrophage differentiation and activation (Bennouna et al., 2003, van Gisbergen et al., 2005). TNF-related ligand B-lymphocyte stimulator (BLyS), also secreted by neutrophils, helps to elicit the proliferation and maturation of B cells (Scapini et al.,

2005). Neutrophil activated pro-chemerin, such as chemerin, can attract both immature DCs and plasmacytoid DCs, and monocytes (Chertov et al., 1997, Wittamer et al., 2005). IL-12 activated neutrophils produce IFN- γ that helps to drive T cell differentiation and macrophage activation (Ethuin et al., 2004).

There is a homeostatic feedback loop among neutrophils, macrophages and T cells regarding neutrophil production in the bone marrow. Granulocyte colony-stimulating factor (G-CSF) is a necessary regulator of neutrophil proliferation and production. Its secretion is regulated by IL-17 produced by regulatory T cells. Extravascular macrophages release IL-23 as the major factor governing IL-17 production. However, apoptosis of neutrophils by macrophage ingestion can suppress IL-23 production.

Many acute and chronic lung diseases including asthma, COPD, bronchiectasis, cystic fibrosis, acute bronchitis, and lung cancer are associated with neutrophilic inflammation. There is abundant evidence showing neutrophils as primary effector cells in COPD (Barnes, 2007). Studies on BAL, sputum, bronchial epithelium and lamina propria indicated that smokers with or without COPD have increased numbers of

activated neutrophils compared with non-smokers (Dragonieri et al., 2009, Janoff et al., 1983, Keatings et al., 1996, Lacoste et al., 1993, Pesci et al., 1998). However, neutrophil numbers are only higher in sputum, BAL and small airways than lung parenchyma large airways (Battaglia et al., 2007), which indicate rapid transit through the airways and parenchyma. Also, the percentage of sputum neutrophils was increased with GOLD stage in COPD patients (Singh et al., 2010).

After maturation in the bone marrow, neutrophils are released into the circulation. When lung inflammation occurs, neutrophils migrate from the pulmonary circulation to postcapillary venules and adhere to endothelial cells in the alveolar wall before passing into the alveolar space (Hogg and Walker, 1995). The recruitment of neutrophils from bronchial circulation to the bronchial submucosa is coordinated by the interaction of adhesion molecules on the surface of neutrophils and proteins released by the bronchial endothelial cells. B₂-integrin (CD18) is a neutrophil adhesion molecule that facilitates neutrophil adhesion to endothelium via its endothelial counter receptor intercellular adhesion molecule 1 (ICAM-1) (Diamond et al., 1990, Woolhouse et al., 2005). Increased expression of macrophage antigen (MAC)-1 (CD11b/CD18) has been found in COPD patients (Noguera et

al., 1998). Also, IL-8, produced by neutrophils, monocytes, alveolar macrophages, pulmonary epithelium, smooth muscle airway cells, eosinophils and fibroblasts, can cause rapid mobilization of MAC-1 to the neutrophil surface (Huber et al., 1991). Increased levels of IL-8 have been detected in BAL and sputum of patients with COPD (Donaldson et al., 2009, Pesci et al., 1998). E-selectin, which is essential to slow down the circulating neutrophils to increase neutrophilic uptake, is upregulated on endothelial cells in the airways of COPD patients (Di Stefano et al., 1994).

Trafficking neutrophils into the respiratory tract from postcapillary venules is predominantly regulated by the epithelial cells that form a barrier lining the airway of the lung (Hogg and Walker, 1995). Chemotactic signals attract neutrophils accumulation near mucosal epithelial cells and in the lumen of the airway (Tanino et al., 2002, Traves et al., 2002). These signals including IL-8 (Tanino et al., 2002), TNF- α (Higham et al., 2000), granulocyte-macrophage colony stimulating factor (GM-CSF) (Aaron et al., 2001), leukotriene B₄ (LTB₄) (Crooks et al., 2000), nuclear factor- κ B (NF- κ B) (Di Stefano et al., 2002), and ENA-78 (epithelial neutrophils activating protein of 78 kDa) (Papi et al., 2006b) have all been found increased in COPD airways. Above chemotactic factors

may derive from alveolar macrophages and epithelial cells, however, IL-8 and leukotriene B₄ (LTB₄) are also produced by neutrophils themselves (Profita et al., 2005). Recently, Blidberg and colleague have shown that chemotaxis of circulating neutrophils towards IL-8 and partly towards LTB₄ was increased in both smokers with or without COPD (Blidberg et al., 2012).

In the respiratory tract of COPD patients, recruited neutrophils have been shown to infiltrate the airway epithelium and submucosal glands from the proximal to the distal part (Saetta et al., 1997), causing these cells to act as producers of potent secretagogues that result in mucus hypersecretion and emphysema (Damiano et al., 1986, Nadel, 1991). Also, increased numbers of neutrophils and CD8⁺ T cells infiltrate the smooth muscle in the peripheral airways of COPD patients compared to smokers and non-smokers, which affects the structure and contractility of airway smooth muscle, contributing to peripheral airways obstruction (Baraldo et al., 2004).

Activated neutrophils produce serine proteases including neutrophil elastase (NE), cathepsin G, proteinase-3, macrophage-derived matrix metalloproteinase (MMP)-8, MMP-

9 and myeloperoxidase (MPO) (Borregaard et al., 2007), which promote the bronchial inflammation, alveolar destruction and also potent mucus stimulants (Ilumets et al., 2008, Weiss, 1989). NE has been demonstrated in emphysematous tissue of COPD patients and there is a correlation between emphysema severity and elastase levels in peripheral blood neutrophils (Damiano et al., 1986, Galdston et al., 1977). Higher concentration of NE-specific fibrinogen degradation product $A\alpha\text{-Val}^{360}$ has been observed in sputum in exacerbations of COPD compared to stable COPD patients (Carter et al., 2012). MMP-9 and MMP-12, which are secreted by macrophages, determine the neutrophils influx (Russell and Ley, 2002). They are also increased in COPD patients (Russell et al., 2002).

Once neutrophils arrive in airways, they generate great amount of O_2^- and reactive oxygen species (ROS), which are also produced by activated neutrophils, eosinophils, macrophages and epithelial cells in the airways of COPD patients (MacNee, 2001b). ROS can damage the epithelium, reduce ciliary beat frequency, stimulate mucus production by goblet cells, and increase the permeability of the bronchial mucosa (Rusznak et al., 1996). All of them lead to airway edema and protein exudation in the airway (Gompertz et al.,

2001). Studies on systemic neutrophils have shown that there is an increase in ROS production in COPD patients compared to healthy smokers and healthy nonsmokers (Gustafsson et al., 2000, Rahman et al., 1996).

Apoptosis of neutrophils is the way to terminate inflammation and further prevent any permanent damage. During apoptosis, neutrophils lose their ability to produce granule contents and undertake phagocytosis (Haslett, 1999). Mucociliary escalator can also remove activated neutrophils if apoptosis is not efficiently performed. However, this removal leads to the production of proteases and chemoattractants that continuously destroy the lung tissue (Nathan, 2006). Increased levels of chemotactic factors, such as IL-8, LTB₄ and GM-CSF, have shown their capacities of delaying neutrophils apoptosis in COPD patients, which leads to secondary necrosis and increases inflammation (Haslett, 1999). Also, delayed apoptosis by neutrophils from COPD patients is related to the expression of bak, bcl-xl and mcl-1 mRNA (Zhang et al., 2012). Interestingly, reduced spontaneous apoptosis only occurs at exacerbation but not stable COPD patients (Pletz et al., 2004).

In contrast to T lymphocytes, particularly CD8⁺ T cells, and macrophages that are predominant inflammatory cells in the lung of patients with mild COPD, neutrophils is more potent in severe COPD patients (Di Stefano et al., 2004). Studies on BAL and sputum from COPD patients indicate that the concentration of neutrophils is positively correlated with airflow limitation (Di Stefano et al., 1998, Stanescu et al., 1996). It has also been found that there are increased numbers of neutrophils in bronchial biopsy specimens and peripheral blood during COPD exacerbations (Papi et al., 2006a, Saetta et al., 1994, Zhu et al., 2001). Furthermore, the percentage of apoptotic blood neutrophils at COPD exacerbations is reduced (Schmidt-Ioanas et al., 2006).

To sum up, neutrophils play an important role in the pathology in COPD, especially at exacerbations. Modulation of neutrophil recruitment and apoptosis in the lungs of patients with COPD may provide potential targets of this disease.

1.2.1.2 Dendritic Cells (DCs)

DCs are potent antigen-presenting cells (APCs) and play a crucial role in the initiation of adaptive immune responses (Banchereau et al., 2000). The first defined pulmonary DCs were Langerhans cells (LCs). They are now recognized by their

immunohistochemical expression of CD1a, langerin (CD207) and by electron microscopy the presence of Birbeck granules (Sertl et al., 1986). Recently, three phenotypically and functionally different pulmonary DC subsets have been classified including type 1 myeloid DCs (mDCs, BDCA1, CD1c⁺), type 2 myeloid DCs (mDCs BDCA3, CD141⁺) and plasmacytoid DCs (pDCs, BDCA2, CD303⁺/CD123⁺). Myeloid and plasmacytoid DCs perform different functions in both innate and adaptive immunity. They have functional plasticity to induce one of three T-cell responses depending on the type of stimuli, including Th1, Th2, or regulatory T (Treg) cells (Vermaelen and Pauwels, 2005).

The role of DCs in the pathogenesis of asthma has been well studied (Lambrecht and Hammad, 2003). However, studies on the involvement of DCs in the development of smokers with COPD are fairly rare. *In vitro* human studies indicate that DC maturation, their capacity to secrete cytokines and expression of adhesion molecules are affected by nicotine. *Mahyar and his colleagues* have shown that in a nicotinic environment (200ug/ml), both endocytic and phagocytic abilities of monocyte-derived DCs are reduced. Nicotine has no effects on DC maturation in response to bacterial antigen lipopolysaccharide (LPS). Nevertheless, they secrete

dramatically lower levels of pro-inflammatory cytokines, such as IL-12, IL-10, and TNF- α . IL-12 is a principal cytokine that bridges innate and specific immune response by activation of NK and T cells and biases the differentiation of CD4 T cells towards IFN- γ -producing Th1 cells (Macatonia et al., 1995). Low level production of IL-12 by nicotine treated DCs indicates that in a nicotinic environment, the ability of DCs to prime type 1 T-cell polarization is diminished. Moreover, T cells co-cultured with DCs in the presence of nicotine produced significantly lower levels of IFN- γ than in the absence of nicotine (Nouri-Shirazi and Guinet, 2003). Another study performed by *Alexandra et al* has shown that nicotine increases the DC expression of adhesion molecules such as the β 2-intergrin lymphocyte function-associated antigen 1 (LFA-1, CD11aCD18) and its ligand CD54 (ICAM-1). Furthermore, the expression of HLA-DR and the DC maturation marker CD83 are also increased in the presence of nicotine (Aicher et al., 2003).

In vivo human studies, the number of CD1a⁺ immature DCs were increased in the alveoli and in the BAL of smokers compared to non-smokers (Casolaro et al., 1988, Soler et al., 1989). However, unlike the airway epithelium that contains exclusive CD1a⁺ DCs, there are normally relatively few CD1a⁺

DCs in human alveoli and BAL (Demedts et al., 2005). In COPD patients, the number of langerin (CD207⁺) immature LCs in the epithelium and the adventitia of small airways is significantly higher compared to never smokers and asymptomatic smokers. This accumulation of DCs is positively correlated to the disease severity. Nevertheless, there is no significant difference in DC numbers in the epithelial layers between current smokers with COPD and ex-smokers with COPD. Also, there is no difference in DC numbers in the lamina propria among groups (Demedts et al., 2007).. Recently, abnormal numbers of DCs subsets were detected in PBMCs from COPD patients. The absolute number of pDCs was significantly reduced along with a marked increase of the mDC/pDC ratio in COPD patients compared to non-smokers (Malta et al., 2013).

CCL20 (also known as MIP3 α) is a chemokine that attracts DCs toward sites of inflammation via interaction with its receptor CCR6 (Reibman et al., 2003). Its mRNA expression is significantly higher in lung tissue from COPD patients compared to healthy non-smokers and healthy smokers. There is an inverse correlation between the FEV₁% and CCL20 mRNA expression. Nevertheless, there is no significant difference in CCR6 mRNA expression in lung tissue between controls and

COPD patients. Similar results have also been found in sputum samples (Demedts et al., 2007). In large airways, current smoking appears to markedly reduce the number of DCs both in the bronchial mucosa and in induced sputum (Reibman et al., 2003). Expression of receptors for antigen recognition, for example, BDCA-1 or Langerin, are significantly increased in mDCs from current smokers with COPD but also markedly decreased in mDCs from ex-smokers with COPD, compared with non-smokers (Stoll et al., 2014).

Overall, investigations on DCs in the pathogenesis of COPD indicate that chronic exposure to cigarette smoking impairs the normal maturation process of DCs and subsequently alters their normal function and interaction with naive lymphocytes, leading to an imbalance of immunity that may increase susceptibility of patients with COPD to respiratory infections.

1.2.1.3 Macrophages

Monocytes, DCs and macrophages, along with neutrophils and mast cells, are all phagocytic cells that are strategically located throughout the body's tissues, where they ingest and process foreign molecules, apoptose cells and debris, maintaining homeostasis, as well as producing multiple mediators to recruit other immune cells (Erwig and Henson,

2008, Mantovani et al., 1972). Macrophages account for the majority of inflammatory cells recovered by airway lavage, regardless of subject status, such as whether or not they are smokers or suffering from airway diseases (Kuschner et al., 1996, Linden et al., 1993). Studies on macrophages have shown that macrophages play a potent role in the pathophysiology of COPD (Barnes, 2004a, Shapiro, 1999).

Numbers of macrophages in BAL fluid, sputum, small airway epithelium, lung parenchyma, bronchial submucosal and bronchial glands are all increased in COPD patients (Barnes, 2004d, Saetta et al., 1993, Saetta et al., 1997, Wallace et al., 1992). In particular, compared to healthy smokers, the number of macrophages is significantly increased by up to 25-fold in the alveolar spaces and tissue in patients with emphysema (Wallace et al., 1992). Also, there is a positive correlation between macrophage numbers in the alveolar walls and the presence of mild-to-moderate emphysema as well as the degree of small airways disease in COPD patients (Di Stefano et al., 1998, Finkelstein et al., 1995, Meshi et al., 2002).

Increased recruitment of monocytes from the circulation in response to monocyte-selective chemokines may be one of

the reasons to account for the increased numbers of macrophages in the lungs of smokers and patients with COPD. Expression of C-C chemokine receptor 2 (CCR2) is thought to be particularly important for macrophage differentiation and recruitment to the airways (Lee et al., 1999), which is 1.4-fold higher levels in macrophages in COPD patients compared to subjects without COPD. Macrophages have a very low proliferation rate in the lungs, however, increased proliferation has been reported to be responsible for the increased numbers of macrophages detected in smokers (de Boer et al., 2000, Barbers et al., 1991, Bitterman et al., 1984). The life span of tissue macrophages ranges from 6 to 16 days. Tissue macrophages are renewed by local proliferation of progenitor cells rather than by monocyte influx into tissue (Sawyer, 1986, Tarling et al., 1987). This feature protects patients from an overwhelming risk of fatal infection when neutrophil production is transiently interrupted. Particles from cigarette smoke can be found within the cytoplasm of alveolar macrophages in smokers for more than two years after smoking cessation (Marques et al., 1997). Therefore, smoking may contribute to the prolonged cellular activation of macrophages. In smokers and patients with COPD, markedly increased expression of the anti-apoptotic protein Bcl-X_L and expression of cyclin-dependent kinase inhibitor p21^{CIP/WAF1} in

the cytoplasm are detected in macrophages (Tomita et al., 2002). These findings suggest that apoptosis of alveolar macrophages may be defective in smokers and in COPD patients, which may be another explanation for the accumulation of macrophages in the lung. In COPD patients and healthy smokers, the expression of apoptosis inhibitor of macrophages in alveolar macrophages is significantly increased in the lung of COPD patients and healthy smokers. Likewise, apoptosis inhibitor of macrophages expression was enhanced in response to CSE exposure (Kojima et al., 2013).

Macrophages tend to display polarized phenotypes by which they can be divided into two major subpopulations, namely, M1 and M2. M1 are classically activated macrophages that exhibit pro-inflammatory properties. These cells play important effector functions including the production of reactive nitrogen species and they secrete high amounts of pro-inflammatory cytokines, such as $\text{TNF-}\alpha$, IL-12 and IL-1 β that regulate antigen presentation to T cells and promote Th1 immunity. In contrast, M2 macrophages have poor antigen presenting capacity but a better clearance of apoptotic cells. They produce anti-inflammatory cytokines, such as IL-10 and TGF- β . Mannose receptor (CD206) and the scavenger receptor CD163 are two representative markers to distinguish M1 and

M2. *In vitro*, GM-CSF can generate M1 macrophages whereas M-CSF can activate M2 macrophages from human peripheral blood monocytes (Murray and Wynn, 2011, Verreck et al., 2006). In the lung, alveolar macrophages predominantly show anti-inflammatory M2 features that have highly phagocytic capacity (Blumenthal et al., 2001, Van den Heuvel et al., 1999). Figure 1.2 shows the macrophage polarization and their features.

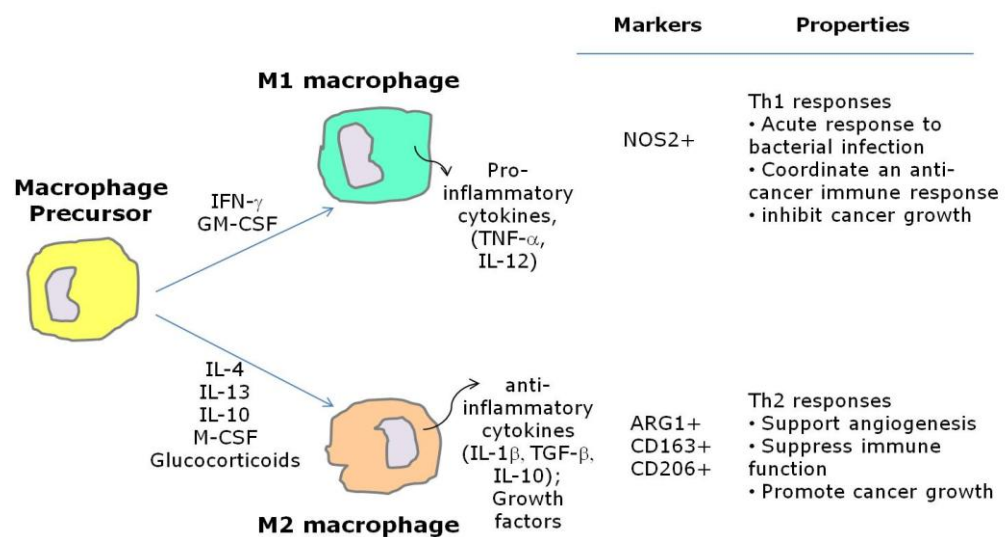


Figure 1.2 Macrophage polarizations. Diversity and plasticity are two hallmarks of macrophages. M1 macrophages (classically activated macrophages) are pro-inflammatory and have a central role in host defense against infection, while M2 macrophages (alternatively activated macrophages) are associated with responses to anti-inflammatory reactions and tissue remodeling, and they represent two terminals of the full spectrum of macrophage activation. *Adapted from: M.C. Schmid & J.A. Varner, J Oncol. 2010:201026*

Decreased phagocytosis of alveolar macrophages has been detected in smokers with COPD and smoking cessation contributes to improvement (Berenson et al., 2006, Hodge et al., 2007, Taylor et al., 2010). This finding suggests that there may be a regulation of macrophage polarization in COPD patients. Kunz and colleagues recently showed that the percentage of macrophages with M2 characteristics is markedly higher in BAL from ex-smokers with COPD compared to current smoking COPD patients. Also, the percentage of M2 in BAL is higher than in induced sputum, which indicates that the M2 macrophage subpopulation is mainly present in the periphery of the lung. In addition, there is no significant difference in the percentage and number of M2 macrophages in BAL between ex-smokers and current smokers suffering from COPD. Although there are other studies to show the shift of macrophage polarization in COPD (Kunz et al., 2011, Frankenberger et al., 2004, Hogger et al., 1998, Kollert et al., 2009), disease severity may be a possible explanation for these controversial findings. Impaired phagocytosis of alveolar macrophages for *Haemophilus influenzae* and *Moraxella catarrhalis* has been detected to be correlated with the severity of COPD (FEV1% predicted) in COPD patients (Berenson et al., 2013). Moreover, Metcalfe and colleague showed that cigarette smoke extract significantly reduced

innate immune response of alveolar macrophages from COPD patients (Metcalfe et al., 2014).

Macrophages induce mucus hypersecretion and lung emphysema via production of LTB₄, IL-1 and matrix metalloprotease enzymes (MMPS). *In vitro*, alveolar macrophages from COPD patients express increased levels of MMP-1 and MMP-9 (Finlay et al., 1997a, Segura-Valdez et al., 2000), and, *in vivo*, expression of MMP-2, MMP-9 and MMP-12 are enhanced in COPD patients (Molet et al., 2005, Montano et al., 2004).

1.2.1.4 T cells

T cells in the lung make up a significant proportion of the total T cell population of the body (Pabst and Tschernig, 1995). They are derived from the progenitor cells that begin recombination of four gene segments, namely, α and β , or γ and δ , which build up the T cell receptor (TCR) genes (La Motte-Mohs et al., 2005, Lauritsen et al., 2006). The TCR is composed of two disulfide-linked chains, either $\alpha\beta$ or $\gamma\delta$. TCR- α/β cells comprise 95% of the T cell population and regulate the immune response by production of cytokines and chemokines. TCR- γ/δ cells play key functions in mucosal

homeostasis and in response to tissue damage at the epithelial surface (Jameson et al., 2003).

α/β T cells

Cytotoxic T cells (Tc cells, CD8⁺) and helper T cells (Th cells, CD4⁺) are α/β T cell subsets. Tc cells consist of 30-40% of the total α/β T cell population and recognize intracellular antigens, such as viral or tumour proteins, presented by MHC class I molecules. Th cells constitute 60-70% of the total α/β T cells and recognize extracellular antigens, such as bacterial peptides, presented by MHC class II molecules (Alam and Gorska, 2003, Parkin and Cohen, 2001). The predominant function of Tc cells is to kill infected cells via producing cytokines such as IFN- γ and toxic molecules like perforin and granzymes (Henkart, 1985, Zagury, 1982). Perforin is a pore-forming protein that helps granule proteins of Tc cells to be transported into target cells (Froelich et al., 1996, Shresta et al., 1999). Granzymes and T-cell intra-cytoplasmic antigen-1 (TLA-1) are two main Tc granule proteins that have the capacity to activate caspase mediated DNA fragmentation resulting in apoptosis of the target cells (Kojima et al., 1994). In addition, binding of Fas-ligand presented by Tc cells to Fas expressed by target cells can also induce apoptosis of infected cells (Nagata and Golstein, 1995). Th cells are coordinating

cells of the immune response. They play a major part in activation of B cells.

Tc cells include two subsets, namely, Tc1 cells and Tc2 cells. Tc1 cells produce IFN- γ whereas Tc2 cells release IL-4 and IL-5 (Mosmann et al., 1997). Th cells are also divided into two main groups, namely, Th1 cells and Th2 cells. Th1 subset secretes IL-2 and IFN- γ and is associated with cell-mediated functions, such as activating Tc cells and delayed hypersensitivity. Although both Th1 and Th2 subset can activate B cells and induce antibody production, the main difference is in the profile of isotype switching- Th1 induce mainly IgG1-3; Th2 induce IgG1-4, IgG3 and IgG A (Mosmann et al., 1997).

Besides CD8⁺ cytotoxic T cells and CD4⁺ helper T cells, Treg cells are a special subpopulation of T cells with the major function of suppressing activation of the immune system by producing suppressor molecules, such as IL-10 or TGF- β (Cederbom et al., 2000, Liu et al., 2003).. CD4⁺ cells expressing high levels of CD25 (CD25 bright) are thought to be Treg cells that represent between 1 and 3% of total CD4⁺ T cells (Dejaco et al., 2006). Besides CD4 and CD25, FoxP3 is mostly restricted to CD4⁺ Treg cells, although some CD8⁺ T

cells do express it (Fontenot and Rudensky, 2004, Tang and Bluestone, 2008). Recently, CD127 has been identified as a unique cell surface marker for Treg cells (Liu et al., 2006, Shen et al., 2009). In particular, CD127^{dim} has been shown to be a specific marker for Treg cells, whereas CD127^{bright} is expressed by non-regulatory T cells.

γ/δ T cells

γ/δ T cells are preferentially associated with epithelial tissue. They are abundant in the gastrointestinal mucosa and skin, which have a potent role in epithelial repair. Unlike α/β T cells, γ/δ T cells do not recognize antigen as peptides presented by MHC molecules. They directly distinguish a great amount of diverse antigens, such as the non-classical MHC class I molecules T10 and T22, without clonal expression (Hayday and Tigelaar, 2003, Holtmeier and Kabelitz, 2005).

In 1995, Finkelstein *et al* showed an increase of CD3⁺ T cells in the lung parenchyma in smokers and there was a positive correlation between the number of T cells of the lung tissue and the extent of emphysema (Finkelstein et al., 1995). Further studies have also shown that numbers of CD4⁺ T cells and CD8⁺ T cells are both increased in the lung parenchyma and airways in COPD patients compared to asymptomatic

smokers and healthy non-smokers, with a predominance of CD8⁺ T cells. This is in contrast to the findings in asthma, in which the predominantly increased cells are CD4⁺ T cells and eosinophils. Abnormal numbers of CD8⁺ T cells in smokers with COPD is not tissue specific. They are detectable in different tissue compartments, such as large airways sub-epithelium (Lams et al., 2000, O'Shaughnessy et al., 1997), peripheral airway (Saetta et al., 1998), parenchyma (Saetta et al., 1999), smooth muscle (Baraldo et al., 2004), bronchial arteries (Peinado et al., 1999, Saetta et al., 1999, Turato et al., 2002), peripheral blood (Urbanowicz et al., 2009), sputum samples (Urbanowicz et al., 2010), and small airways such as BAL (Saetta et al., 2003, Saetta et al., 2002). Some studies demonstrate the correlation between the number of CD8⁺ T cells and lung function (Di Stefano et al., 2001, O'Shaughnessy et al., 1997, Turato et al., 2002). However, recently one study demonstrated that the distribution of T-cell subsets in BAL of patients with mild to moderate COPD is smoking *pe se* but not correlated with airway obstruction (Forsslund et al., 2014). Smokers with or without COPD show higher percentages of CD8⁺ T cells and NKT-like cells than those never smokers and ex-smokers with COPD. Although toll-like receptors (TLRs) are expressed at low levels in the lung tissue, the percentage of CD8⁺ T cells expressing TLRs

including TLR1, 2, 4, 6 and 2/1 is significantly elevated in smokers with COPD compared to smokers with normal lung functions (Freeman et al., 2013). Activated CD8⁺ T cells in COPD secrete Th1 predominant cytokine pattern that includes increased production of IFN- γ , interferon-inducible protein-10 (IP-10), and monokine induced by interferon-gamma (MIG). In turn, these mediators can cause tissue destruction through the up-regulation of MMP production by macrophages and other immune cells (Grumelli et al., 2004, Maeno et al., 2007). Enzymes released by CD8⁺ T cells, such as granzymes and perforin, as well as binding of Fas-ligand to Fas between CD8⁺ T cells and target cells, can also cause cell apoptosis (Henkart, 1994, Kojima et al., 1994).

Although studies of CD4⁺ T cells are less extensive than CD8⁺ T cell infiltrates, CD4⁺ T cells have also been shown to be increased in airways, lung parenchyma and peripheral blood in COPD patients (O'Shaughnessy et al., 1997, Saetta et al., 1999), particularly in proximity to bronchus associated lymphoid tissue (Hogg et al., 2004). Decreased number of CD4⁺ helper T cells is detectable in BAL from smokers with or without COPD in comparison with non-smoking participants (Forsslund et al., 2014). CD4⁺ helper T cells are mainly responsible for orchestrating downstream immune process by

producing cytokines. They are essential for the full development of adaptive immune cytotoxicity by priming and promoting the long-term survival of CD8⁺ T cells. They also help activation and differentiation of antibody elaborating B-cells (Wan, 2010).

Treg cells (CD4⁺CD25⁺ T cells) are a subpopulation of CD4⁺ T lymphocytes with significant anti-inflammatory and immunomodulatory effects (Bluestone and Tang, 2005, Jiang and Chess, 2006). Recently, more studies have shown interests in the Treg cells in the pathogenesis of COPD. However, the findings are conflicting. Smyth *et al* demonstrated the up-regulation of Treg cells in BAL fluid in healthy smokers and COPD patients, compared to healthy non-smokers, and there is a positive correlation between FoxP3 expression and the number of pack-years (Smyth *et al.*, 2007). Nevertheless, Barcelo *et al* found increased levels of Treg cells only in BAL from smokers but not in COPD patients and healthy non-smoking participants. Also, the number of Treg cells were higher in BAL than in the peripheral blood in the three groups of individuals studied (Barcelo *et al.*, 2008). Human Treg cells are classified into three subsets including CD25(++)CD45RA(+) resting Tregs (rTregs), CD25(+++)CD45RA(-) activated Tregs (aTregs), which play

suppressive functions, and CD25(++)CD45RA(-) cytokine secreting (Fr III) cells. Decreased proportions of rTregs and aTregs, and increased proportion of F γ III cells among CD4⁺ T cells were found in COPD compared to normal smokers (Hou et al., 2013).

γ/δ T cells play a potent role in tissue repair and mucosal homeostasis. However, only a few studies address the potential role of γ/δ T cells in COPD. Richmond et al showed the number of γ/δ T cells was increased in the bronchial glands of healthy smokers compared to healthy non-smokers (Richmond et al., 1993). Likewise Majo et al also found increased γ/δ T cells in the lung parenchyma of healthy smokers (Majo et al., 2001). Recently, two studies both confirmed that the number of γ/δ T cells are also increased in both peripheral blood and BAL fluid from healthy smokers, compared to COPD patients and healthy non-smokers, particularly in BAL (Ekberg-Jansson et al., 2000, Pons et al., 2005). Also, current smokers with normal lung function showed a significant up-regulation of γ/δ T cells compared to ex-smokers with normal lung function (Pons et al., 2005). Importantly, the active smoking inducing up-regulation of γ/δ T cells was blunted in current smokers with COPD, which

indicates the capacity for tissue repair involving γ/δ T cells might be postponed in COPD patients.

Cytokines, chemokines and enzymes producing by T cells perform different roles in cell recruitment, cell activation, cell differentiation as well as killing infected cells. Molecular mechanisms of these molecules in COPD will be demonstrated individually in later sections.

1.2.1.5 NKT cells

NKT cells share receptor structure and function with both T cells and NK cells (Kronenberg and Gapin, 2002). They express T cell membrane proteins, such as CD3, CD4 and CD8, as well as NK cell markers including CD56, CD161 and inhibitory natural killer cell receptors (Emoto and Kaufmann, 2003). The most widely studied subset of NKT cells is invariant NKT (iNKT), which express both an invariant TCR consisting of the α -chain ($V\alpha 24/J\alpha Q$ in humans and $V\alpha 14/J\alpha 281$ in mice) paired with a β -chain ($V\beta 11$ in human and $V\beta 2$, $V\beta 7$ or $V\beta 8$ in mice) and NK cells markers, such as NK1.1 and Ly-49 (Berzins et al., 2004, Stenstrom et al., 2004). The unique invariant TCR of NKT cells recognizes glycolipid antigens, such as the synthetic glycolipids α -galactosylceramide (α -Galcer) and the endogenous

isoglobotrihexosylceramide (iGb3) by interaction with the MHC class I-like molecule CD1d presented by antigen presenting cells (Burdin and Kronenberg, 1999, Crowe et al., 2003, Zhou et al., 2004). However, recognition of CD1d by NKT cells appears to be tissue specific (Hong et al., 1999). α -Galcers has stronger immunostimulatory activities and has been used as a powerful tool for analyzing NKT cell function *in vivo* because conventional T cells do not recognize this molecule (Crowe et al., 2003). There is an overwhelming body of evidence showing that most NKT cells are thymus-dependent, however, some argue for an extrathymic origin (Bendelac, 1995, Bendelac et al., 1997, Crowe et al., 2003, MacDonald, 1995).

In humans, most studies on NKT cells have been limited in peripheral blood. However, one study has shown that NKT cells are clearly detectable in human liver at approximately 4% of hepatic T cells (Doherty et al., 1999). Recently, it has been shown that iNKT cells are found in low numbers in the BAL fluid (less than 2% of total CD3⁺ T cells), induced-sputum (less than 1.3% of CD3⁺ T cells) and bronchial biopsy (less than 1.7% of CD3⁺ T cells) in subjects with mild or moderately severe asthma, subjects with stable or exacerbated COPD and healthy volunteers (Vijayanand et al., 2007). Furthermore,

there are no significant differences among those three groups. The distribution of NKT cells in other human tissues remains to be determined.

According to the expression of CD4 or CD8 surface molecules on NKT cells, they are classified into three subpopulations, namely, CD4⁺CD8⁻ NKT cells (CD4 NKT cells), CD4⁻CD8⁺ NKT cells (CD8 NKT cells) and CD4⁻CD8⁻ NKT cells (DN NKT cells). Recent studies have highlighted the distinct Th1- and Th2-type cytokine profiles of these subsets of NKT cells (Gumperz et al., 2002, Takahashi et al., 2002, Takahashi et al., 2000). Upon stimulation, the CD4 NKT cells produce Th1 cytokines (e.g. IFN- γ) and Th2 type cytokines (e.g. IL-4), plus IL-2, IL-10 and granulocyte-macrophage colony-stimulating factor (GM-CSF). By contrast, CD8 NKT cells and DN NKT cells predominantly secrete Th1 cytokines (e.g. IFN- γ) but not IL-2, GM-CSF or Th2 cytokines (Kim et al., 2002, Lee et al., 2002). The different cytokine production of NKT cell subpopulations has different actions on immune effector cells in adaptive immune responses. For example, CD4 NKT cells induce the greatest activation of NK cells, CD4⁺ T cells and B cells but have no effect on CD8⁺ T cells. These observed stimulatory activities might be involved in both Th1- and Th2-type immunity. Compared with the other two NKT cell subsets, CD8

NKT cells have substantially greater stimulatory activities on CD8⁺ T cells while producing less IFN- γ .

One of the interesting features of the NKT cells is its paradoxical function in different diseases including malignancies and autoimmune dysfunctions. It is hard to conclude which kind of NKT cell subset performs more important roles in immune response. This is dependent on the distribution of NKT cell subsets in different tissues. For example, DN NKT cells are at greater frequency than CD4 NKT cells and CD8 NKT cells in human liver (Kenna et al., 2003). Moreover, activation of NKT cells can be either beneficial or detrimental (Duarte et al., 2004, Kikuchi et al., 2001, Kim et al., 2005, Sharif et al., 2002, Terabe et al., 2000). Previous studies have raised the prospect of manipulating NKT cells for therapeutic benefits (Parekh et al., 2004, Yu et al., 2005). Phase I cancer clinical trials revealed that soluble α -GalCer treatment is safe, but exerts moderate immunostimulatory effects (Major et al., 2004, Nakai et al., 2004, Tupin et al., 2004). Depending on differences in cytokine profiles in NKT cell subsets, non-selective activation of all NKT cell subsets could likely result in unexpected immunological outcomes for the intended therapeutic use. This suggests that clinical trials to develop the therapeutic potential of NKT cells should

probably utilize strategies that activate or bias toward specific NKT cell subpopulations rather than the whole NKT cell population. For examples, the numbers of NKT cells were reduced significantly in smokers with COPD compared with non-smoking healthy volunteers and healthy smokers whereas the relative proportion of CD8⁺ NKT cells was increase in peripheral blood from COPD patients (Urbanowicz et al., 2009).

In addition, cytokine release by NKT cells is always rapid and transient, with a peak of activity within 24 hours. After 24 hours, the production of these cytokines, for example IFN- γ , quickly returns to baseline levels (Lin et al., 2006). Thus, the timing of activation of NKT cells is also an important factor to achieve therapeutic efficacy.

Besides cytokine production, NKT cells also exhibit potent cytotoxic functions. They can induce cell apoptosis by constitutively expressing Fas-L that interacts with Fas⁺ target cells (Arase et al., 1994). They also kill tumor targets in a perforin-dependent manner (Smyth et al., 2000).

The capacity of rapidly producing high levels of immunoregulatory cytokines may allow NKT cells to regulate adaptive immunity and autoimmune diseases. Previous studies

have shown that NKT cells are critical in both innate and adaptive immune responses against viral and bacterial infections (Grubor-Bauk et al., 2003, Hansen and Schofield, 2004, Kakimi et al., 2000, Skold and Behar, 2003), as well as in tumour immunity and certain autoimmune and allergic diseases such as type 1 diabetes and allergic asthma (Lisbonne et al., 2003, Smyth et al., 2000).

Recently, Kim *et al* introduced a new mouse model to detect invariant NKT cell function in both asthma and COPD (Kim et al., 2008). They found that, in mouse lung tissue, the chronic pulmonary symptoms required interactions between CD4⁺ NKT cells and macrophages via CD1d molecules. IL-13, derived from iNKT cells, also served as a source of early anti-inflammatory cytokine with great activity on activation of macrophages. This activation resulted in an autocrine signalling loop due to more IL-13 produced by activated macrophages (Martinez et al., 2009). Similar results were seen in the COPD group. Increased numbers of macrophages expressing high levels of IL-13 were observed in COPD affected lungs removed from transplant recipients compared with normal lungs (Kim et al., 2008).

To date, a number of studies have shown differences of distribution and function of NKT cells in peripheral blood, induced sputum and tissues. For instance, Urbanowicz *et al* have demonstrated remarkable findings on NKT cell features in peripheral blood from COPD patients (Urbanowicz *et al.*, 2009). The numbers of NKT cells were reduced significantly in smokers with COPD compared with non-smoking healthy volunteers and healthy smokers whereas the relative proportion of CD8⁺ NKT cells was increased. The expression of cytotoxic effector molecules, namely, perforin and granzyme B, were both markedly lower in COPD subjects. In contrast to smoking-free healthy participants and healthy smokers, NKT cells isolated from peripheral blood of COPD patients also showed less cytotoxic activity. Su *et al* further confirmed and extend these findings (Chi *et al.*, 2011). The number of iNKT cells was significantly reduced in peripheral blood from COPD patients compared to healthy smokers. Furthermore, the frequency of iNKT cells is lower in peripheral blood from patients with exacerbations of COPD compared to stable COPD patients. Interestingly, the proportion of NKT cells is dramatically higher in induced sputum from smokers with COPD compared to healthy smokers and healthy non-smokers (Urbanowicz *et al.*, 2010). Also, the expression of both perforin and granzyme B by NKT cells is also increased in

COPD patients compared to the other two groups, which means that NKT cells from COPD patients are more cytotoxic than those from healthy smokers and non-smoking subjects.

NKT cells, as a unique lymphocyte population sharing characteristics with both conventional T cells and NK cells, are involved in immediate immune responses through their efficient cytokine secretion and lytic activity. Therefore, it is necessary to investigate their roles in health and disease. Previous studies on NKT cell performance in COPD highlight that it is important to design clinical studies aiming to modify NKT cell activity for therapeutic benefit.

1.2.1.6 Natural killer (NK) cells

NK cells, as a critical innate immune effector cell type, are an important source of immunoregulatory cytokines and play potent roles in early host defence by interaction with other immune cells. They are equipped with a great number of receptors that can be either inhibitory or stimulatory and also possess certain adhesion molecules that facilitate NK cell homing and trafficking (Gregoire et al., 2007, Lanier, 1998, Lanier et al., 1998). Upon activation, NK cells release a number of cytokines including myeloid-differentiation or activation factors (IL-3, TNF- α , GM-CSF), IFN- γ and type 2

cytokines (IL-4, IL-5, IL-10 and IL-13) and cytotoxic molecules, such as granzymes and perforin (Cooper et al., 2001a, Cooper et al., 2001b, Loza et al., 2002). These important properties warrant NK cells multiple functions in both innate and adaptive immune system. Killing infected cells by cytotoxic molecules, inducing adaptive immunity through cytokine and chemokine secretion, and providing co-stimulatory molecules to T cells and B cells are main immunological functions of NK cells.

CD56 is an isoform of the human neural cell adhesion molecule and widely used as a cell surface marker to identify NK cells (CD3⁻CD56⁺). Although its function in NK cells is still unexplored, early studies suggest that CD56 may function as an adhesion protein mediating interactions between NK cells and target cells (Lanier et al., 1989, Nitta et al., 1989, Suzuki et al., 1991). CD16 (FcγRIII), as a receptor of IgG, is also detectable on NK cells (Dall'Ozzo et al., 2004, Lanier et al., 1986). According to the intensity of CD56 and CD16 expression, NK cells have been classified into two subsets including CD56^{bright} and CD56^{dim} NK cells (Cooper et al., 2001a). In peripheral blood, CD56^{bright} NK cells that rarely express CD16 molecule only comprise of approximately 10% of total NK cells and function as cytokine-producing cells with

low level expression of perforin (Konjevic et al., 1995, Nagler et al., 1989). CD56^{dim} NK cells expressing high density of CD16, however, are a majority (~90%) of NK cells and defined as killers due to their high cytotoxicity. The perforin levels of these cells are also high (Nagler et al., 1989). Besides CD56 and CD16, cytokine receptors, such as receptors for IL-2 and IL-15, and other activation or inhibitory receptors are also expressed on NK cells. They can selectively recognize MHC class I ligands, MHC class-I related ligands and host-encoded non-MHC ligands (Lanier, 1998).

Adhesion molecules are thought to play functions in cell-cell interactions, trafficking and homing. Two subpopulations of NK cells have distinct expression patterns of these molecules that result in their functional diversity. In contrast to CD56^{dim} NK cells that express high levels of LFA-1 and PEN5 (a sulfated polylactosamine carbohydrate epitope), CD56^{bright} NK cells highly express CC-chemokine receptor 7 (CCR7) and L-selectin that both are important for homing immune cells to secondary lymphoid organs (Frey et al., 1998), which also may be a reason to explain why CD56^{bright} NK cells are highly presented in lymph node in contrast to CD56^{dim} NK cells (Ferlazzo et al., 2004). Figure 1.3 presents the phenotypes of NK cells with their functional molecules.

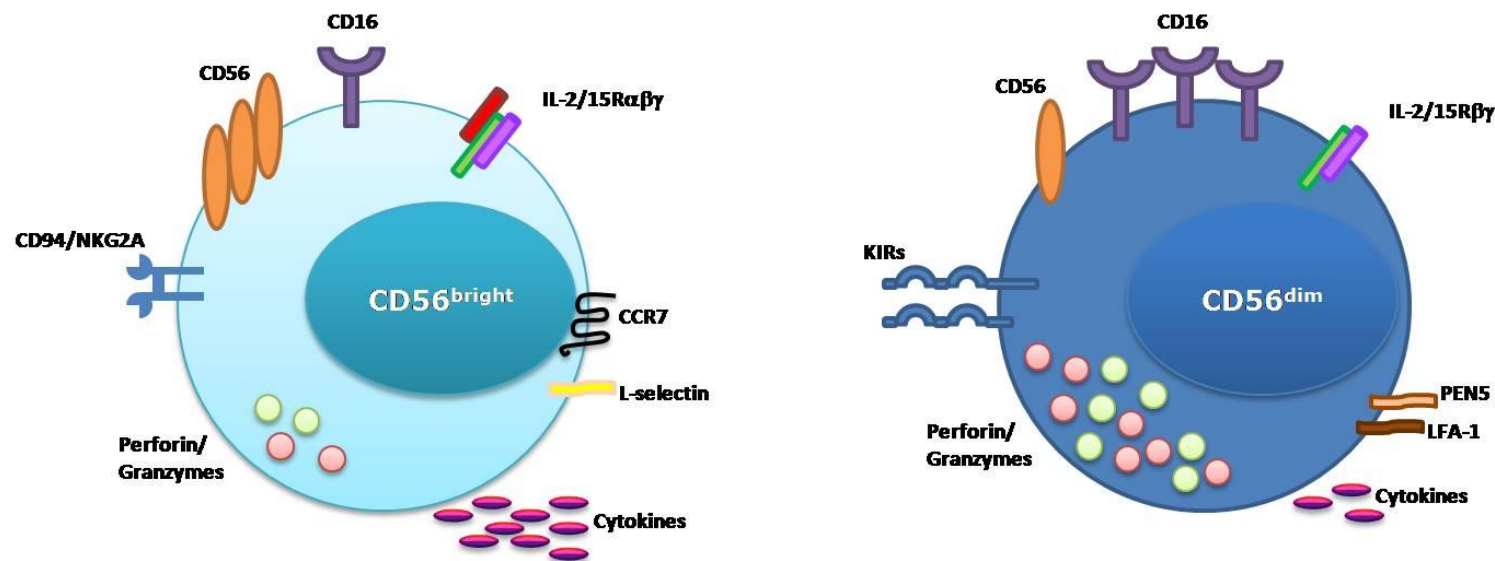


Figure 1.3 Properties of human NK cell subsets. CD56^{bright} NK cells (left in the figure) express high levels of inhibitory receptor CD94/NKG2A, cytokine receptors IL-2/15Rαβγ and adhesion molecules including L-selectin and CCR7. However, CD16 and KIRs are rarely expressed on these cells. Upon activation, these cells can release high levels of cytokines. Thus, the predominant function of CD56^{bright} NK cells is immunoregulatory. By contrast, CD56^{dim} NK cells express high levels of CD16 and KIRs. They can secrete a great amount of cytotoxic proteins including perforin and granzyme B but release few cytokines after stimulation. Therefore, they mainly perform as killer cells in the immune system. (Adapted from Megan et al. The biology of human natural kill-cell subsets. 2001. *TRENDS in Immunology*. 22: 633-640)

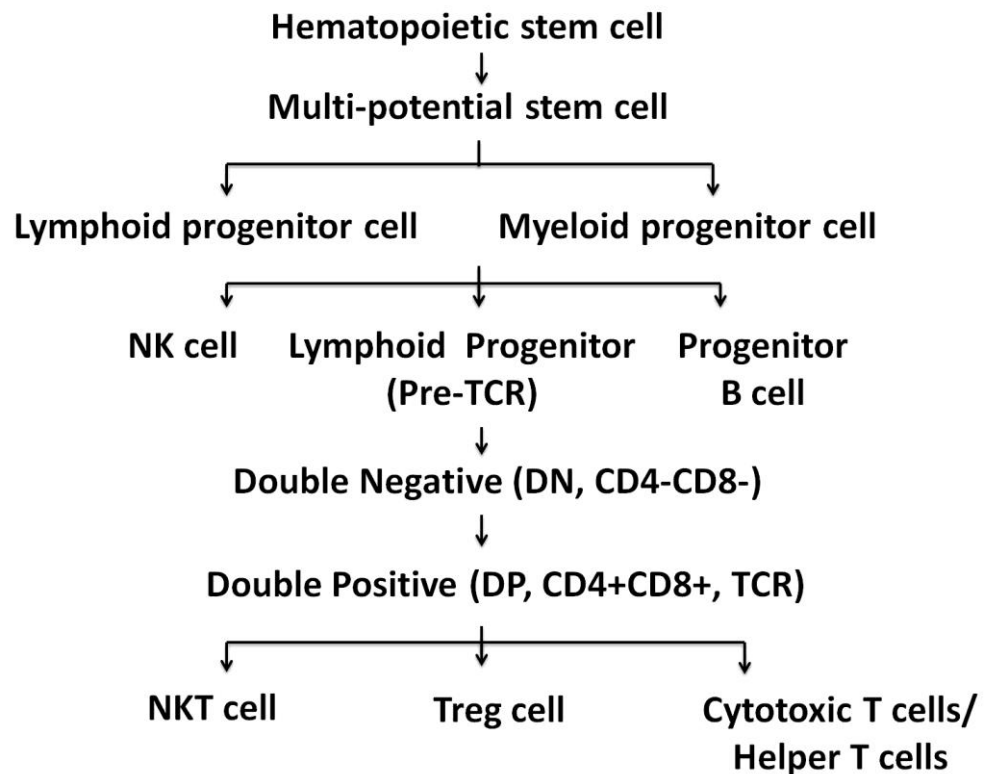


Figure 1.4 Origins of NK and NKT cells. Hematopoietic stem cells give rise to all blood lineages. NK cells are innate immune effector cells. They are derived from either lymphoid or myeloid progenitor cells, and CD3-CD56+. NKT cells are a distinct lineage of T lymphocytes that are derived from lymphoid progenitor, and CD3+CD56+.

T cells, B cells and NK cells all develop from haematopoietic progenitor cells that reside in the CD34⁺ cell compartment (Lotzova et al., 1993). Figure 1.4 shows the Origins of NK and NKT-like cells in hematopoiesis. Recent studies indicate that bone marrow and lymph nodes are two major physiological sites for NK cell development. IL-2 has been detected as an

effective promoter to support the differentiation of NK cells from CD34⁺CD38⁻ hematopoietic progenitors (Lotzova et al., 1993, Shibuya et al., 1995). Exogenous IL-2 can also expand NK cell proportions (Caligiuri et al., 1993, Piguet et al., 1986, Rosenberg et al., 1987). IL-15 is another necessary promoter for NK cell development. It has structural similarities with IL-2 and they both share several functions, such as inducing T cell proliferation, generation and activation of NK cells. Studies focusing on the NK cell progenitor in bone marrow have shown that IL-15 plays a critical roles in both the development and the maturation of NK cells (Fehniger and Caligiuri, 2001, Williams et al., 1999). It can induce human CD34⁺ haematopoietic progenitor cell differentiation into NK cells in the absence of other cytokines (Mrozek et al., 1996). However, differentiation based on IL-15 is not able to enhance NK cell expansion (Mrozek et al., 1996), unless two additional bone marrow stromal factors, namely, ligands of the receptor tyrosine kinases c-kit and flt-3, act synergistically with IL-15 (Lyman and Jacobsen, 1998, Yu et al., 1998). Besides bone marrow, lymph node is another important site for NK cell development. CD56^{bright} NK cells have been found in peripheral blood when CD34⁺ progenitor cells are treated with recombinant IL-15 (Lyman and Jacobsen, 1998). Although CD56^{bright} NK cells only comprise around 10% of circulating NK

cells, they are a majority of NK cells (~90%) in lymph nodes. Interestingly, in lymph node, addition of IL-2 can functionally differentiate CD56^{bright} NK cells to CD56^{dim} NK cells through increasing their cytotoxicity (Ferlazzo et al., 2004). Collectively, lymph nodes may serve as a harbour for NK precursor cells that more frequently develop into CD56^{bright} NK cells.

NK cell receptors

NK cell receptors can be classified into three families including receptors recognizing MHC class I ligands, receptors interacting with MHC class I related ligands, and receptors for host-encoded non-MHC ligands. NK receptors for MHC class I refer to those with capacities to recognize either classical or non-classical class I molecules encoded by genes within the MHC. In human two representative receptors in this family have been investigated, namely, human killer cell immunoglobulin-like receptors (KIRs) and conserved CD94/NKG2 receptors. These receptors are suggested to play predominant functions in monitoring cells for aberrant expression of MHC class I molecules in immune surveillance.

KIRs, belonging to the Ig superfamily, are also referred to as the CD158 family. They are type I transmembrane glycoproteins that contain two (D1 and D2) or three (D0, D1

and D2) Ig-like domains (designated KIR2D and KIR3D respectively) in the extracellular region (Colonna and Samaridis, 1995, Wagtmann et al., 1995), a short stalk region separating the Ig-like domains from the transmembrane segment, and cytoplasmic domains which characteristics are variable in length, for instance, some receptors possess long (L) cytoplasmic domains containing at least one intracellular immunotyrosine-based inhibitory motif (ITIM) sequence, they are named as KIR2DL or KIR3DL. While other receptors with short (S) cytoplasmic domains possess no ITIM sequence, which are called KIR2DS or KIR3DS. There are also five different types of cytoplasmic tails in the structure of KIRs. Numbers (1-5) designating to which cytoplasmic tail group each KIR belongs are also used to label KIRs, such as KIR2DL1, KIR2DS2 and so on. In humans, 14 different KIR genes have been detected, they are KIR2DL (1-5), KIR2DS (1-5), KIR3DL (1-3) and KIR3DS1 (Raulet et al., 2001). Unlike the ITIM-induced signalling pathway in KIR2DL and KIR3DL, KIR2DS and KIR3DS lacking ITIMs possess a Lys residue in their transmembrane domains. This charged residue associates with an immunoreceptor tyrosine-based activation motif (ITAM)-bearing DAP12 adapter protein to transmit signals. In other words, KIRs with long cytoplasmic domains perform as inhibitory receptors, whereas KIRs possessing

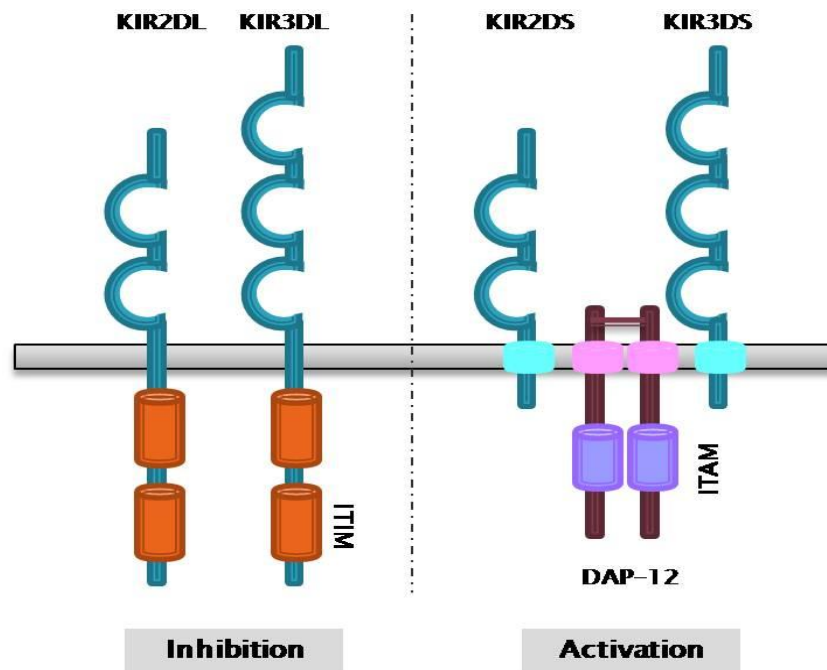


Figure 1.5 Representative structures of KIRs (KIR2DL, KIR3DL, KIR2DS and KIR3DS). KIRs contain long (L) cytoplasmic domains possessing at least one ITIM sequence. Thus their major function is inhibitory. Whereas KIRs with short (S) cytoplasmic domains lack ITIMs, they possess a Lys residue in their transmembrane domains. This charged residue associates with an ITAM-bearing DAP12 adapter protein to transmit signals. KIR(S) function as activating receptors. (Adapted from Jordan et al. Natural killer cells in human disease. 2006. Clinical Immunology. 118: 1-10)

short cytoplasmic domains are assumed to be activating receptors. Figure 1.5 shows representative structures of KIRs.

In humans, ligands for KIRs are mainly HLA subfamily, such as HLA-A, HLA-B and HLA-C. KIR2DL can recognize HLA-C

proteins (Colonna et al., 1992), whereas KIR3DL shows the capacity of ligation with HLA-B and certain HLA-A (Gumperz et al., 1997, Gumperz et al., 1995). To date, ligands for KIR2DS and KIR3DS are not known; all studies show that these activation KIRs either do not interact with HLA class I molecules or bind them with much weaker affinity than that of the paired inhibitory KIRs (Valiante et al., 1997, Winter et al., 1998).

KIR2DL4 is a unique gene in KIR family with remarkable structural features. HLA-G is suggested to be KIR2DL ligand in some studies (Boyson et al., 2002, Cantoni et al., 1998, Rajagopalan and Long, 1999).

Most human KIR genes are expressed by subsets of NK cells, memory and effector CD8⁺ T cells, some CD4⁺ T cells and $\gamma\delta$ TCR⁺ T cells except KIR2DL4 (Ferrini et al., 1994, Phillips et al., 1995). Once they are expressed, they are stably maintained on the cell surface. However, KIR2DL4 again as an exception – it is expressed constitutively on the surface of CD56^{bright} NK cells and on only activated CD56^{dim} NK cells (Goodridge et al., 2003).

Interaction of inhibitory KIRs with their MHC class I ligands expressed on target cells can lead to suppression of cytokine production and cytotoxicity of KIR-bearing NK cells and T cells. Although several studies have speculated that lacking an inhibitory receptor for self MHC class I molecule can elicit autoimmune disorders (Namekawa et al., 2000, Snyder et al., 2003, Suzuki et al., 2004, Yen et al., 2001), which is detrimental, KIRs can also be beneficial in immune responses to tumours or viral pathogens (Flores-Villanueva et al., 2001).

CD94/NKG2 receptors are type II transmembrane proteins of the C-type lectin family. They contain two C-lectin domains in the extracellular region, a transmembrane region and a cytoplasmic domain (Lazetic et al., 1996). In humans a single CD94 gene is genetically linked to four NKG2 family genes (NKG2A-, -C, -E, and -F) (Glienke et al., 1998, Houchins et al., 1991). CD94 does not bind ligands of CD94/NKG2 complex and also plays no function on signal transduction (Lazetic et al., 1996, Lohwasser et al., 1999). Unlike KIRs, in humans, CD94/NKG2 receptors recognize non-conventional MHC class I molecules (HLA-E) (Braud et al., 1998).

Although NKG2A and NKG2C share the same ligand HLA-E, their immunological functions are different. Vales-Gomez *et al*

has found that inhibitory receptor CD94/NKG2A shows stronger affinity than the activating receptor CD94/NKG2C. In addition, peptides bound to HLA-E can also affect binding selection between the two receptors (Llano et al., 1998, Vales-Gomez et al., 1999). To date, ligands and functions of CD94/NKG2E and CD94/NKG2F have not been well characterised.

Nearly same as KIRs, CD94/NKG2 receptors are also expressed on most NK cells, memory and effector CD8⁺ T cells and $\gamma\delta$ TCR⁺ T cells. However, their expression on CD4⁺ T cells is undetectable (Lazetic et al., 1996). Also, CD94/NKG2 expression is not constitutively stable and is regulated by surrounding cytokines, such as IL-12, IL-15 and TGF- β (Bertone et al., 1999, Derre et al., 2002, Mingari et al., 1998).

Besides KIRs and CD94/NKG2 receptors, leukocyte immunoglobulin (Ig)-like receptors (LILRs) are another type of receptors for MHC class I molecules. One representative member of LILRs family is LILRB1 that has the inhibitory activity. LILRB1 are expressed on a subset of T cells, B cells and monocytes. They can also be found on NK cells from peripheral blood with variable frequency (undetectable to 75%) (Colonna et al., 1997, Cosman et al., 1997). However, its

affinity of ligation with all HLA molecules seems very low (Chapman et al., 1999, Lepin et al., 2000). U18 has been investigated as a ligand with higher affinity to LILRB1 and can elicit their inhibitory activity during viral infection (Beck and Barrell, 1988, Cosman et al., 1997, Chapman et al., 1999).

There is a kind of glycoprotein which is not encoded by genes in the MHC complex but structurally related to MHC class I molecule. It does not serve as a peptide-binding molecule to present antigens to T cells. However, this MHC class I-related protein can be recognized by NK receptor NKG2D (Raulet, 2003). NKG2D, as a type II transmembrane-anchored glycoprotein, is a unique activating receptor which both structurally and functionally differ from other NKG2 receptors (Bauer et al., 1999, Jamieson et al., 2002). It is encoded by a single gene without polymorphism and does not need to form dimers with CD94 (Wu et al., 1999). In human, NKG2D is expressed on the surface of NK cells, γ/δ TCR⁺ T cells, and especially, it is constitutively present on all CD8⁺ T cells (Wu et al., 1999). The expression of NKG2D can be regulated by cytokines. For example, IL-15 and TNF- α can enhance the NKG2D expression whereas TGF- β causes downregulated effects on it (Castriconi et al., 2003, Lee et al., 2004). Unlike two isoforms of NKG2D (NKG2D-L and NKG2D-S, "L" and "S"

represent long or short protein respectively) found in mouse, humans only express NKG2D-L that transmit signals through transmembrane segment associating with adaptor protein DAP10 (Diefenbach et al., 2002, Wu et al., 1999). Studies demonstrate that in the absence of DAP10, NKG2D is not able to be presented on cell surface and only retained in the cytoplasm (Wu et al., 1999). Human ligands for NKG2D have been identified, such as MICA, MICB, ULBP1, ULBP2, ULBP3 and ULBP4 (Bauer et al., 1999, Cosman et al., 2001). Figure 1.6 presents the interactions of the NKG2D and CD94 family with target cells.

Besides receptors for recognition of MHC class I and MHC class I-like molecules, NK cells also possess additional receptors that can encounter targets lacking expression of MHC class I or MHC class I-related proteins. However, this type of receptors is supposed to be costimulatory receptors because they are unable to individually initiate an immune response. In human, this kind of receptors mainly includes 2B4 receptor (CD244), DNAM-1 receptor (CD226) and CD16 (Fc γ RIII).

2B4 (CD244) and its ligand CD48 are both members of the CD2 family of Ig-related proteins. In human, they are expressed by all NK cells, most γ/δ TCR⁺ T cells, memory and

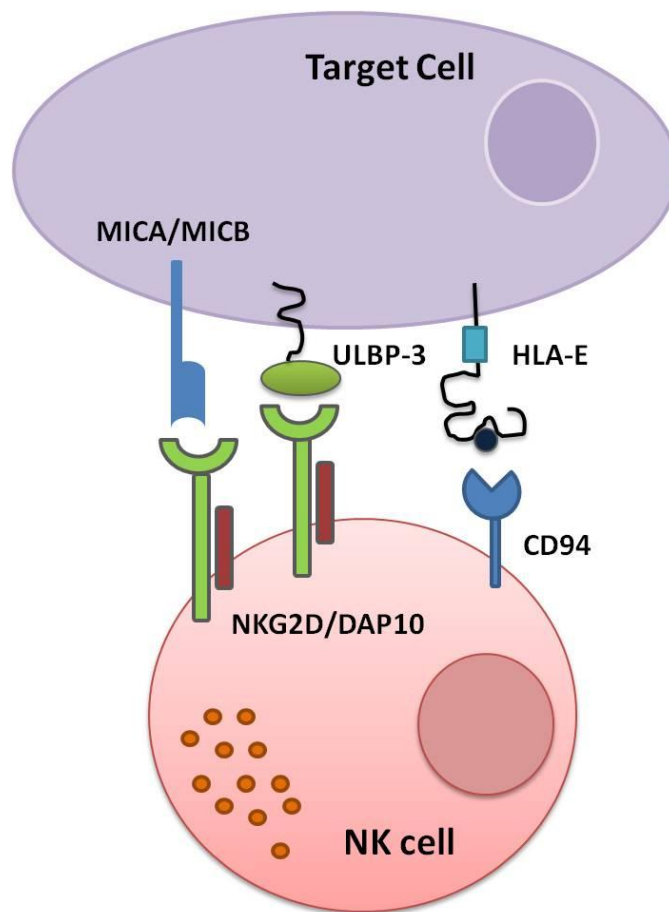


Figure 1.6 Interactions of NKG2D and CD94 family with target cells. CD94 receptors are type II transmembrane proteins of the C-type lectin family. In humans a single CD94 gene is genetically linked to four NKG2 family genes (NKG2A, -C, -E and -F). CD94 receptors recognize non-conventional MHC class I molecules (HLA-E). NKG2D is a type II transmembrane-anchored glycoprotein. Human only expresses NKG2D-L that transmits signals through transmembrane segment associating with adaptor protein DAP10. Human ligands for NKG2D are MICA, MICB, ULBP1-4.

effector CD8⁺ T cells, monocytes and basophils (Nakajima et al., 1999, Speiser et al., 2001, Tangye et al., 2000). Although signalling pathways of 2B4 receptor is still unresolved, studies

in human suggest that 2B4 may serve as a multifunctional receptor that can cause either inhibitory or activating responses depending on the stage of NK cell differentiation and activation (Namekawa et al., 2000, Parolini et al., 2000).

DNAX accessory molecule-1 (DNAM-1, also called CD226) is a member of the Ig superfamily and ligated with CD112 and CD155 that are broadly distributed on many tissues (Tahara-Hanaoka et al., 2004). Expression of DNAM-1 is detectable in human NK cells, a majority of T cells, a subset of B cells, monocytes and platelets (Castriconi et al., 2004, Pende et al., 2006, Reymond et al., 2004, Scott et al., 1989, Shibuya et al., 1996).

CD16 is a low affinity receptor for IgG. It binds the lower hinge region of IgG and induces activation signals through Fc ϵ RI γ -chain and CD3 ζ , which enable NK cells to mediate antibody-dependent cellular cytotoxicity (ADCC) (Dall'Ozzo et al., 2004, Radaev et al., 2001). Expressions of ICAM-1 together with IgG on infected cells are required to induce efficient ADCC (Bryceson et al., 2005). CD16 shows high expression on CD56^{dim} cells.

In conclusion, since NK cells are defined as non-MHC-restricted cells according to their capacities of recognizing target cells that either express or lack MHC class I molecules, missing-self hypothesis demonstrates that self-MHC molecules engages inhibitory receptors on the surface of NK cells, which can prevent attack against normal cells (Ljunggren and Karre, 1990). Actually, rather than absolutely saying that interaction between MHC-I molecules and inhibitory receptors are terminated, responses of NK cells to target cells are regulated by a balance of signals from inhibitory and activating receptors (Figure 1.7). Either lack of the ligands for inhibitory receptors or signals from activating receptors that are stronger than inhibitory signals can both induce NK cell stimulation. On the other hand, NK cells display their inhibitory functions if there are no activating ligands presented in target cells or much stronger inhibitory signals are available. Several experimental lines of evidence have confirmed this corollary. When activating receptors are sufficiently stimulated or multiple activating receptors are engaged, NK cells still can eliminate infected cells, although MHC class I molecules on these infected cells are still ligated with inhibitory receptors (Cerwenka et al., 2001, Diefenbach et al., 2001, Lanier et al., 1997). Also, Fc-receptors (FcRs) have similar scenarios in immune response. By virtue of coexpression of activating and

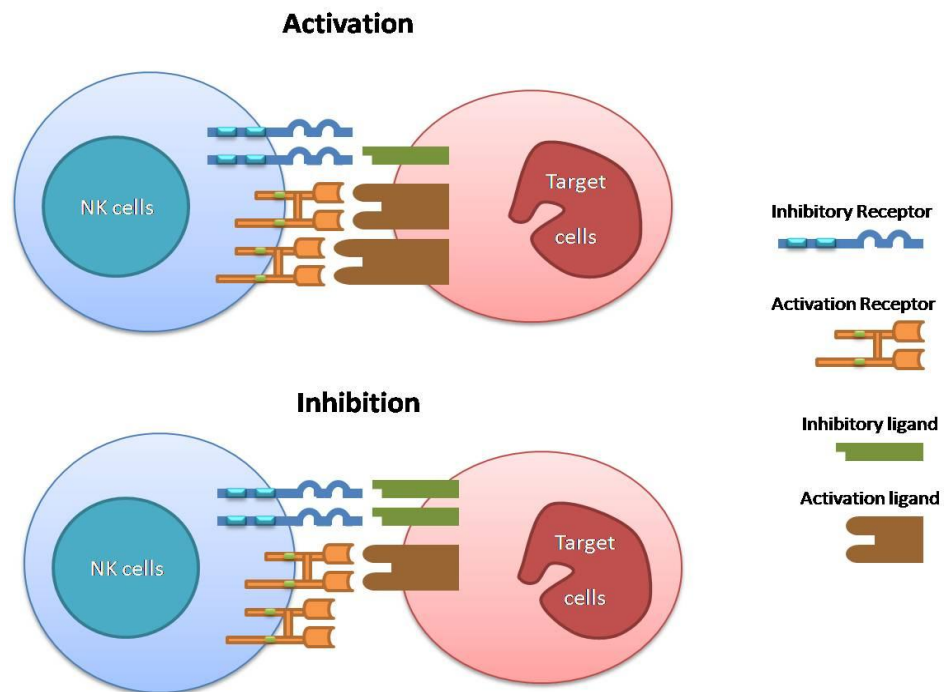


Figure 1.7 Regulation of NK cell responses by inhibitory or activating receptors. NK cell activation can be induced in two manners. Either lack of the ligands for inhibitory receptors or signals from activating receptors that are stronger than inhibitory signals can both induce NK cell stimulation. On the other hand, NK cells display their inhibitory functions if there are no activating ligands presented in target cells or much stronger inhibitory signals are available. (Adapted from Lewis L. Lanier. NK cell recognition. 2005. Annu. Rev. Immunol. 23: 225-274)

inhibitory signalling FcRs on the same cells, they set a threshold for immune cell activation by immune complexes. Loss of the balanced signalling leads to uncontrolled response that can result in the damage of healthy tissues and ultimately to the initiation of autoimmune processes (Abes et al., 2009).

So far, several major types of NK cell receptors presented in human cells have been introduced with respect to their ligands specificity and signalling properties. However, other receptors expressed on NK cells, such as natural cytotoxicity receptors and cytokine receptors, are also important for generating various NK cell functions in response to target ligands. Although many features and functions of NK cells have been well studied in tumour, viral infection and autoimmune diseases, their immunological properties in COPD patients are still unexplored. Since the predominant risk factor of COPD is cigarette smoking, several studies have successfully detected the effects of cigarette smoke in humans. *In vivo*, previous studies have shown that NK cell numbers and their phagocytic activities are suppressed in both asymptomatic smokers and COPD patients (Ferson et al., 1979, Prieto et al., 2001). Our group further demonstrated that decreased numbers of NK cells are present in peripheral blood from COPD patients compared to healthy non-smokers and healthy smokers

(Urbanowicz et al., 2009). Also, their cytotoxic function is defective. Nevertheless, NK cell proportions and their cytotoxicity are increased in induced sputum from COPD patients in contrast to healthy non-smokers and healthy smokers (Urbanowicz et al., 2010). In addition, cigarette smoke treated NK cells from non-smoking healthy subjects show defective cytotoxicity and their capacity for cytokine secretion is also impaired upon poly I:C activation (Mian et al., 2008). In conclusion, investigation of NK cells' properties in COPD may provide new insights for understanding the development of COPD as well as indicate more beneficial therapeutic approaches for the disease.

1.2.1.7 Epithelial cells

The surface of airway epithelium consists of ciliated cells, goblet cells, ciliated cells and basal cells. These cells work together as an efficient barrier against pathogens through intercellular epithelial junctions. They also exert important roles in airway defence mechanisms, such as clearance of the mucus, water transport, antioxidant, antiprotease functions and cell apoptosis (Puchelle et al., 2006).

Cigarette smoking is the key risk factor in the development of COPD. Injury in airway epithelium caused by cigarette

smoking can result in increased proliferation of airway epithelial cells, which further lead to the squamous metaplasia. Likewise, squamous metaplasia can decrease mucociliary clearance and contribute to the increased risk of squamous cell carcinoma in COPD patients (Cosio et al., 1978, Papi et al., 2004).

A series of studies have shown that cigarette smoking can not only cause the increased expression of IL-8 and mucin in epithelial cells but also reduce the function of trans-epithelial electric resistance (TEER), which reflects a breakdown of the epithelial integrity that can affect the defence mechanisms of the airway (Glader et al., 2006a, Rusznak et al., 1999, Richter et al., 2002, Takeyama et al., 2001). Also, up-regulation of early growth response gene 1 (Egr-1) was detectable when epithelial cells were exposed to cigarette smoke. Egr-1 mediates the production of pro-inflammatory cytokines, such as IL-1 β , TNF- α , which have shown increased levels in smoking conditions (Reynolds et al., 2006). Furthermore, *in vitro*, epithelial cells from small airways of healthy smokers and COPD patients secrete more TGF- β than do those from healthy non-smokers (Takizawa et al., 2001). Increased levels of fibroblast growth factors (FGF) 1 and 2 are also observed in

the bronchial epithelium of COPD patients (Kranenburg et al., 2005).

Pathogens including bacteria and viruses can exacerbate COPD and accelerate deterioration of lung function (Rohde et al., 2003, Seemungal et al., 2001). Previous studies have shown that rhinovirus (RV) is the dominant viral pathogen in exacerbation of COPD. RV infection can cause secretion of a variety of pro-inflammatory chemokines, such as ENA-78, IL-8 and IP-10, as well as cytokines, for example, IL-1, IL-6, IL-11, IL-16, TNF- α and GM-CSF, by epithelial cells (Proud and Chow, 2006). These inflammatory mediators play different functions on recruitment and activation of inflammatory cells including neutrophils, macrophages, T cells and NK cells that can induce pulmonary and systemic inflammatory responses, mucus hypersecretion and airway remodelling, which result in the exacerbation of COPD. Beside viruses, bacteria can also cause exacerbation of COPD by colonizing and infecting the lower respiratory tract (Murphy, 2006). Although *M. catarrhalis* and H. influenza can trigger immune response in airway epithelial cells (Shuto et al., 2001, Slevogt et al., 2006, Watanabe et al., 2004), a study performed by Sethi *et al.* has shown that alteration in bacterial load was not an important mechanism for exacerbation of COPD (Sethi et al., 2007).

Airway epithelial cells are the initial targets of reactive oxygen species (ROS). Pierrou *et al.* have investigated extensive changes in oxidant response gene expression in the airway epithelial cells that are related to cigarette smoking and the presence of COPD (Pierrou *et al.*, 2007). Furthermore, the levels of 4-hydroxy-2-nonenal-modified (4-HNE) proteins were increased in airway epithelial cells of COPD patients (Rahman *et al.*, 2002). 4-HNE is a diffusible and highly reactive lipid peroxidation end-product. The modification of 4-HNE can lead to abnormal regulation of cell proliferation and inhibition, T cells apoptosis and activation of signalling pathways.

Hypersecretion of mucus and reduction of mucociliary clearance are two main mechanisms inducing mucus accumulation (Danahay and Jackson, 2005, Stannard and O'Callaghan, 2006). The epithelial goblet cells are a main source of secretory gel-forming mucins in the airway. The number of goblet cells has shown increased level in COPD patients compared to healthy smokers (Innes *et al.*, 2006). Mucociliary clearance is dependent on the interactions between the ciliated epithelium, the height of the periciliary fluid and mucus (Stannard and O'Callaghan, 2006). It has been shown that there is significant reduction (nearly no presence) of ciliated cells in COPD patients, compared to

healthy non-smokers and healthy smokers (Rusznak et al., 2000).

Apoptosis is a potent mechanism for elimination of damaged or infected cells. There are abundant studies showing that numbers of apoptotic cells, including bronchial and alveolar epithelial cells and endothelial cells, are increased in the parenchyma in COPD patients (Henson et al., 2006, Hodge et al., 2005b, Hodge et al., 2003a, Park et al., 2007). The apoptosis of airway epithelial cells is disease related rather than smoking status. Hodge S et al. have shown that there is no significant difference in apoptosis of airway epithelial cells between ex-smokers with COPD and current smoking COPD patients (Hodge et al., 2005b), which implies that apoptosis exists after smoking cessation.

1.2.1.8 Mast cells

Since neutrophils, macrophages and CD8⁺ T cells are commonly considered the predominant cells in the pathogenesis of COPD (Chung and Adcock, 2008), the role of mast cells in COPD is just beginning to become unravelled.

Mast cells are highly heterogeneous cells and originate from myeloid stem cells (Anderson et al., 1990). Their maturation

and differentiation are influenced by many cytokines and chemokines, such as IL-3, 4, 9 and 10 (Brown et al., 2008). They are normally classified into two subsets, namely, mucosal mast cells (MC_T) and connective tissue mast cells (MC_{TC}). The MC_T mainly exist in the respiratory tract and gut, and their number can be modulated in different inflammatory conditions, such as allergy and inflammatory bowel disease. The MC_{TC} are predominantly present in connective tissue, such as the skin, and their number is constant (Irani et al., 1986).

Studies related to the mast cell functions in lung diseases are mostly performed in animal models (Galli et al., 2005, Sekizawa et al., 1989, Thomas et al., 1992). However, recently studies have shown the number, the density and mediators released by mast cells are altered in airways of smokers, even in COPD patients. Wen *et al* have explored that the numbers of mast cells were significantly increased in sputum of smokers compared to ex-smokers (Wen et al., 2010). In the lung and skin of smokers, there is also an increase in absolute numbers of mast cells (Kalenderian et al., 1988). Furthermore, elevated histamine and tryptase levels in BAL fluid from smokers have also been demonstrated (Bessa et al., 2008, Kalenderian et al., 1988). Gosman *et al* have investigated the distribution and the numbers of tryptase- and

chymase- positive mast cells in the airways (Gosman et al., 2008). There was no significant difference in the distribution of these cells in the airways between COPD patients and healthy volunteers. In central airways, the concentration of these mast cells was significantly higher in the subepithelial area, whereas in the peripheral airways, the highest concentration of these cells existed in the adventitia. However, numbers of these mast cells differed between COPD patients and controls. The number of chymase- positive mast cells was significantly higher in both subepithelial areas and smooth muscle of peripheral airways in COPD patients than in controls. Whereas the number of tryptase- positive mast cells in the subepithelial area of central airways was lower in COPD patients than in controls. Moreover, higher percentages of both tryptase- and chymase- positive mast cells were detected in the peripheral airways and positively correlated with FEV₁% predicted, which was compatible with the issue that peripheral airways are the main site of airflow limitation in COPD (Hogg et al., 2004, Hogg et al., 1968).

Besides population and distribution, the density of mast cells in the lung of COPD patients has been investigated lately (Hogg et al., 1968). In the small airways, there were significant decreases in total mast cell density in both patients

with severe COPD and patients with moderate to severe COPD compared to healthy non-smokers and healthy smokers. In the pulmonary vessel walls, the reduction of mast cell density was only shown in severe COPD subjects but not other groups. In addition, the balance of MCTC and MCT was shifted. The population of MCT was decreased in all anatomical compartments of the lung, whereas the density of MCTC was markedly higher in the walls of small airways and in alveolar parenchyma in severe COPD groups than other groups. Recently, Soltani and colleague have shown that in COPD airways, both reticular basement membrane and lamina propria mast cells were increased (Soltani et al., 2012)

Taken together, although alternation of mast cell numbers, density and distribution has been identified in several studies, the mechanisms about the signalling pathway and by which mast cells can be activated in the airways of COPD patients are still not well explored. Mast cells express many receptors, such as abundant TLRs, and can release leukotrienes, cytokines and chemokines after activation (Kumar and Sharma, 2010). Therefore, evaluation of expression of key mast cell associated molecules can contribute to more understanding of the role of mast cells in COPD.

1.2.1.9 Eosinophils

Eosinophils are end-stage cells derived from the bone marrow. Their maturation and proliferation are regulated by GM-CSF, IL-3 and IL-5 (Denburg, 1999). Eosinophil cationic protein (ECP) and eosinophil peroxidase (EPO) are two eosinophil specific proteins. They are toxic to bronchial epithelial cells (Blanchard and Rothenberg, 2009). Eosinophils, together with mast cells and basophils, are the predominant source of cysteinyl-leukotrienes (Bozza et al., 1997). Eosinophilic inflammation is mostly studied in allergic airway inflammation (Walsh et al., 2010). Recently, studies on its role in COPD have started to be investigated. Some groups have demonstrated increased numbers of inactive eosinophils in the airways and lavage of stable COPD patients, whereas others have failed to present differences in eosinophil numbers in airway biopsies, BAL, or induced sputum (Lacoste et al., 1993, Turato et al., 2001). Interestingly, the levels of ECP is higher in induced sputum from moderate to severe COPD in the absence of eosinophils (Brightling et al., 2005, Brightling et al., 2000, Gibson et al., 1998, Takahashi et al., 2006), which indicates that eosinophils may have degranulated during disease development (Keatings and Barnes, 1997). A negative correlation between FEV₁ and the ratio of activated eosinophils to total eosinophils in endobronchial biopsies has been

reported in COPD patients (Lams et al., 2000). Furthermore, levels of sputum eosinophils and ECP concentration are also negatively correlated with FEV₁ (Balzano et al., 1999). Eosinophil function has been accepted to be important in severe COPD. Greater numbers of eosinophils have been detected in bronchial biopsies and BAL fluid during acute exacerbation of COPD compared with stable COPD and controls (Saetta et al., 1996, Saetta et al., 1994).

1.2.2 Cytokines in COPD

The potential roles of cytokines in COPD have been explored in many studies. Over 50 cytokines have now been identified in COPD (Barnes, 2004d). However, cytokines work in a complex manner. They may have different or even opposing actions in different cell types or in different inflammatory process.

Proinflammatory cytokines in COPD

1.2.2.1 Tumour necrosis factor- α (TNF- α)

TNF- α is a member of peptide ligands that stimulate a corresponding series of structurally related receptors (Bazzoni and Beutler, 1996). It plays important roles in many inflammatory diseases. Proinflammatory cells, such as macrophages, monocytes, dendritic cells, B cells, CD4⁺ T cells, neutrophils, mast cells and eosinophils, and structural cells

including fibroblasts, epithelial cells and smooth muscle cells, are all good sources for TNF- α production (Russo and Polosa, 2005). Increased levels of TNF- α have been found in induced sputum from COPD patients, particularly during exacerbations (Aaron et al., 2001). Soluble TNF- α receptors are also increased in sputum of COPD patients (Vernooy et al., 2002). Serum concentrations of TNF- α are elevated in COPD patients with weight loss, particularly during exacerbations (Calikoglu et al., 2004), which indicates that TNF- α may have functions in the cachexia of severe COPD (de Godoy et al., 1996, Di Francia et al., 1994, Pitsiou et al., 2002). There is a significant increase in TNF- α released from circulating cells in COPD patients compared to normal controls. Plasma concentrations of TNF- α are also increased slightly in COPD patients compared to normal subjects during exercise (Rabinovich et al., 2003).

TNF- α activates NF- κ B pathway and p38 mitogen-activated protein kinase (MAPK) that switch on the transcription of inflammatory genes including cytokines, chemokines and proteases released by epithelial cells and macrophages, which indicates that TNF- α play a role in amplifying the inflammation of COPD, such as activation of neutrophils, monocytes, macrophages, epithelium, mucus secretion and destruction of

lung parenchyma (Aggarwal, 2003). Furthermore, TNF- α accounts for 70% of cigarette smoke-induced emphysema in mice (Churg et al., 2004).

1.2.2.2 Interleukin-1 β (IL-1 β)

IL-1 β is a proinflammatory cytokine with similar functions as TNF- α (Watkins et al., 1999). It markedly stimulates alveolar macrophages to produce inflammatory cytokines, chemokines and MMPs in COPD patients (Culpitt et al., 2003). The concentration of IL-1 β is increased in induced sputum of COPD patients and is correlated with disease severity. However, concentrations of IL-1 receptor antagonist and soluble IL-1 receptor are reduced in same COPD groups (Sapey et al., 2009).

1.2.2.3 IL-6

IL-6 is a marker of inflammation and plays functions on both proinflammatory response and anti-inflammatory response. Its effects are determined by the presence of other cytokines (Barnes et al., 2011). Therefore, it provides a link between innate and acquired immunity. Increased concentrations of IL-6 are detected in induced sputum, exhaled breath and plasma of COPD patients, especially during exacerbation (Bucchioni et al., 2003, Grubek-Jaworska et al., 2012, Hageman et al.,

2003, Wedzicha et al., 2000). Levels of IL-6 released by monocytes are higher in COPD patients compared to normal subjects (Aldonyte et al., 2003).

1.2.2.4 Granulocyte-Macrophage Colony Stimulating Factor (GM-CSF)

GM-CSF is a monomeric protein and is secreted by T cells and macrophages. It is a strong chemoattractant for neutrophils. It also has the ability of enhancing oxidative metabolism and phagocytotic activity of neutrophils and macrophages (Hamilton, 2002). There is a higher concentration of GM-CSF in BAL fluid from stable COPD patients, particularly during exacerbation (Balbi et al., 1997).

1.2.2.5 IL-32

Recently, IL-32, as a newly explored proinflammatory cytokine, has been shown to be induced by IFN- γ . Upon activation of NF- κ B and p38 MAPK, IL-32 stimulates the production of TNF- α and IL-1 β (Joosten et al., 2013). Increased expression of IL-32 has been found in epithelial cells, macrophages and CD8⁺ T cells in COPD patients, and is correlated with disease severity (Calabrese et al., 2008).

1.2.2.6 Thymic stromal lymphopoietin (TSLP)

TSLP is a cytokine belonging to IL-7 family. It regulates the secretion of chemokine ligand by dendritic cells including CCL-17, CCL-22 and CXCL-10, which attract T lymphocytes to the airways (Liu et al., 2007). Increased expression of TSLP has been detected in airway epithelium of COPD patients (Ying et al., 2008). In addition, elevated levels of TSLP can be found in airway smooth muscle of COPD patients (Zhang et al., 2007).

T cell cytokines

1.2.2.7 IL-12 family

IL-12, secreted by macrophages, DCs and airway epithelial cells, plays a potent role in Th1 cell differentiation and activation (Watford et al., 2003). It also up-regulates the IFN- γ production (Trinchieri and Scott, 1995). Its effects are mostly mediated by activation of STAT4. IL-23, shared p40 subunit with IL-12, plays an important role in regulation of Th17 cells (Qu et al., 2013). It also activates STAT4. IL-27 is another member of the IL-12 family (Bosmann and Ward, 2013, Gee et al., 2009). It has effects on Th1 cell differentiation via a STAT1-dependent mechanism. Increased levels of IL-12 have been detected in bronchial biopsies of COPD patients (Di Stefano et al., 2004). There is also elevated

expression of IL-23 in the bronchial mucosa of COPD subjects (Di Stefano et al., 2009b). A study performed by Huang et al has mentioned that polymorphisms of the IL-27 gene are linked to increased risk of COPD (Huang et al., 2008).

1.2.2.8 Th17 cytokines

Th17 cells are a subset of CD4⁺ T cells. Their differentiation is regulated by IL-1 β , IL-6, and IL-23 (Manel et al., 2008). IL-17A, IL-21 and IL-22 are major cytokines secreted by Th17 cells (Chen and O'Shea, 2008). Traves *et al* have shown increased levels of IL-17A in the sputum of COPD patients (Traves and Donnelly, 2008). Furthermore, the expression of IL-17A, IL-22 and IL-23 are all markedly elevated in the bronchial mucosa of stable COPD patients (Di Stefano et al., 2009b). Interestingly, IL-17A levels are also increased in smokers with normal lung function, which indicates that the secretion of IL-17A is mediated by smoking *per se* rather than disease itself. Increased expression of IL-22 is also observed in the bronchial epithelium and submucosa of COPD patients compared to healthy smokers and healthy non-smokers.

1.2.2.9 Other T cell cytokines

IL-2, IL-4, IL-9, IL-13, IL-15, IL-18 and IFN- γ are all produced by either Tc cells or Th cells (Mosmann and Coffman, 1989).

Higher IL-2 levels have been found in lung plasma samples from stable COPD patients compared to participants with rapidly progressive COPD (D'Armiento et al., 2009). Also, corticosteroid (dexamethasone) has shown less inhibition of IL-2 production in COPD patients compared to healthy volunteers although these numerical differences between groups did not reach statistical significance (Kaur et al., 2012).

Levels of both IL-4 released by CD8⁺ T cells and IFN- γ released by CD4⁺ T cells are increased in the BAL fluid from COPD patients (Barczyk et al., 2006). There is also an increase in the expression of IFN- γ in bronchial biopsies of COPD patients (Di Stefano et al., 2004a).

IL-9 is normally released by Th2 cells and has functions in allergic inflammation (Shimbara et al., 2000). It increases the expression of MUC5AC and therefore, is a potent inducer of mucus production (Reader et al., 2003). Interestingly, there is a significant increase in the expression of IL-9 in T lymphocytes from bronchial biopsies of COPD patients (Panzner et al., 2003).

IL-13 is another inducer of mucus production and plays a role in the differentiation of goblet cells through activation of EGFR

(Shim et al., 2001). Although there is no significant difference in the expression of IL-13 between COPD and controls (Grubek-Jaworska et al., 2012), smokers with mucus hypersecretion show a higher concentration of IL-13 in bronchial biopsies compared to non-smokers (Miotto et al., 2003).

IL-15 shares several similar functions with IL-2. Details of their features will be indicated in the later chapter. Recently, *in vitro*, it has been found that IL-15 stimulation by itself had no effect on IFN- γ protein expression in COPD patients at GOLD stages 2, 3, and 4. However, IL-15 primed CD8⁺ lung T cells to have greater cytotoxic potential by expressing higher levels of intracellular perforin when they encounter a target cells in COPD (Freeman et al., 2010). Furthermore, CSE treated NK cells were less activated after poly I:C stimulation, and this suppression was IL-15 mediated (Mian et al., 2009).

IL-18 is not only secreted by T cells but also activated macrophages (Nakanishi et al., 2001). IL-18 protein is highly expressed by alveolar macrophages, CD8⁺ T cells and both the bronchiolar and alveolar epithelia in the lung of COPD patients. Its serum levels are also elevated in smokers with or without COPD compared to non-smokers (Imaoka et al., 2008). IL-18

concentrations are also increased in the induced sputum of COPD patients and correlated with disease severity (Rovina et al., 2009). Higher IL-18 gene expression has been found in plasma and in skeletal muscle of COPD patients (Petersen et al., 2007), which indicates that IL-18 may function in systemic manifestations of COPD.

1.2.2.10 IL-10

IL-10 is an anti-inflammatory cytokine secreted by monocytes, alveolar macrophages and Treg cells. Decreased concentration of IL-10 has been detected in induced sputum of COPD patients (Barnes, 2001). However, its production seems to be elevated in macrophages from normal smokers (Lim et al., 2000). Interestingly, increased expression of IL-10 is found in bronchial biopsies from COPD patients (Panzner et al., 2003).

To sum up, although many cytokines are abnormally expressed in COPD patients, overlapping effects of them mean therapeutic benefits may be trivial. Nevertheless, exploring the cytokine network in COPD provides a new insight for understanding the pathogenesis of disease development.

1.2.3 Chemokines in COPD

COPD involves chronic inflammation of small airways and lung parenchyma, with the recruitment of inflammatory cells. Trafficking inflammatory cells to infected areas is orchestrated by multiple chemokines and cytokines. There are over 50 different chemokines that have been indentified. They can selectively activate 19 different surface receptors expressed by target inflammatory cells (Locati et al., 2005). Unlike classical cytokine receptors, receptors for chemokines belong to the 7 transmembrane receptor super family of G-protein-coupled receptors (GPCRs), which makes it possible to discover small molecule inhibitors (Proudfoot, 2002). Four different families of chemokines are now categorized, namely CC, CXC, Cm and CX3C chemokines, based on their differences in the position of critical cysteine residues. Binding of chemokines to their corresponding receptors results in the activation of signal transduction pathways, which further induces chemotaxis or other cellular activities (Gutierrez-Ramos et al., 2000). Currently, several chemokines and their receptors seem to be involved in the pathogenesis of COPD (de Boer, 2005).

1.2.3.1 Interleukin-8

Interleukin-8 (IL-8, CXCL8) is a CXC chemokine and a predominant chemoattractant of neutrophils (Hammond et al.,

1995) . It activates two chemokine receptors, namely, CXCR1 and CXCR2. CXCR1 is a low affinity receptor that is almost specific for IL-8, whereas CXCR2 is a high affinity receptor and is shared by other related CXC chemokines (Premack and Schall, 1996). Binding of IL-8 to these two receptors can activate protein kinase B and GTPases, which consequencely enhances the neutrophil adherence to endothelial cells and directs cell migration. Activated protein kinase B can also stimulate the phosphoinositide 3 kinase that induces F-actin polymerization, leading to pseudopod formation and chemotaxis (Chodniewicz and Zhelev, 2003). In addition, Ras and MAP kinases in neutrophils are activated when IL-8 binds to its receptors, which induces their degranulation (Bowman et al., 1998).

IL-8 is synthesized in several cell types, predominantly macrophages, epithelial cells and neutrophils (Mukaida, 2003), up on activation by several stimuli including $\text{TNF-}\alpha$, $\text{IL-1}\beta$, bacterial products including LPS, certain viruses, oxidative stress and cigarette smoke extract (DeForge et al., 1993, Johnston et al., 1998, Kwon et al., 1994, Nakamura et al., 1991, Schulz et al., 2004). Release of IL-8 is regulated transcriptionally by several transcription factors, such as $\text{NF-}\kappa\text{B}$, p38MAP kinase and ERK pathways (Vaugh and Wilson,

2008). NF- κ B is the predominant transcription factor for IL-8 secretion (Carter et al., 1998). Therefore, inhibition of the NF- κ B activating kinase IKK2 results in the reduction of IL-8 production (Hacker and Karin, 2006, Tang et al., 2001).

Previous studies have confirmed that IL-8 is a major chemokine that plays key roles in the neutrophilic inflammation in COPD. The concentration of IL-8 is markedly increased and correlated with increased numbers of neutrophils in both induced sputum and BAL fluid from COPD patients (Keatings et al., 1996, Nocker et al., 1996, Soler et al., 1999, Yamamoto et al., 1997). Unsurprisingly, neutrophils themselves release IL-8 that attracts more neutrophils. Therefore, a self-perpetuating inflammation may be established in COPD patients (Bazzoni et al., 1991). In induced sputum, IL-8 levels are elevated during acute exacerbation and are positively correlated with the bacterial colony count, which indicates that bacterial infection may at least in part induce neutrophilic inflammation through induction of IL-8 secretion in the airways (Hill et al., 2000, Patel et al., 2002). In BAL, IL-8 is the only chemokine with higher concentration in smokers with emphysema, compared to healthy smokers (Tanino et al., 2002). Furthermore, IL-8 concentrations are also increased in hospitalized COPD

patients and are correlated with skeletal muscle weakness (Spruit et al., 2003).

More IL-8 production has been observed in cultured airway epithelial cells and alveolar macrophages from COPD patients than smokers with normal lung function, which also indicates an amplified response (Culpitt et al., 2003, Schulz et al., 2004). Alveolar macrophages provide a great source of IL-8 in response to stimuli. Cells derived from COPD patients release more IL-8 than those from health smokers (Culpitt et al., 2003). IL-8 protein and mRNA are increased in bronchiolar epithelial cells from COPD patients (de Boer et al., 2000). Also, there is elevated basal release of IL-8 from airway epithelial cells of COPD patients (Profita et al., 2003, Schulz et al., 2003).

In clinical trials, anti-IL-8 antibodies decreased only approximately 30% of the neutrophil chemotactic activity of sputum from COPD patients (Beeh et al., 2003, Crooks et al., 2000). This is because other neutrophil chemotactic factors, such as LTB₄ and C5a, are also involved. Therefore, blocking IL-8 alone is not sufficient as a therapeutic strategy to reduce neutrophil inflammation in the respiratory tract. CXCR2 is a receptor of IL-8 and is also shared by other chemokines.

Small antagonists of CXCR2 are now in clinical development for the treatment of COPD (Chapman et al., 2009).

1.2.3.2 CXCL1 (growth-related oncogene- α , GRO- α)

CXCL1 is another CXC chemokine that is involved in COPD. It is released by alveolar macrophages and airway epithelial cells upon TNF- α and IL-17 activation (Jones and Chan, 2002, Prause et al., 2003, Schulz et al., 2004). It also binds to CXCR2 expressed by neutrophils and monocytes (Premack and Schall, 1996). CXCL1 concentration is significantly increased in both induced sputum and BAL fluid from COPD patients compared to healthy smokers and healthy non-smokers (Traves et al., 2002). There are greater amounts of CXCL1 released by epithelial cells from COPD patients compared to healthy smokers (Schulz et al., 2004), and by BAL cells from normal smokers compared to non-smokers (Morrison et al., 1998). In COPD patients, monocyte chemotactic activity is increased in response to CXCL1, which may account for the higher amount of alveolar macrophages in the lungs of COPD patients (Traves et al., 2004).

1.2.3.3 CXCL5 (epithelial-cell-derived neutrophil activating peptide-78, ENA-78)

CXCL5 is mainly released by epithelial cells and also binds to CXCR2 (Imaizumi et al., 1997). Unlike IL-8 and CXCL1, CXCL5 does not induce an increased chemotactic activity on monocytes (Traves et al., 2004). The concentration of CXCL5 is increased in the BAL fluid from COPD patients compared to healthy non-smokers (Tanino et al., 2002). However, there is no significant difference between patients with emphysema and healthy smokers. Similarly to CXCL1, the amount of CXCL5 derived from BAL cells is higher in COPD patients compared to healthy non-smokers (Morrison et al., 1998). Furthermore, CXCL5 expression is significantly increased in epithelial cells during exacerbations of COPD (Qiu et al., 2003). Recently, Stefano and colleague showed that the numbers of CXCL5⁺ endothelial cells in the bronchial submucosa is negatively correlated with COPD severity (Di Stefano et al., 2009a).

1.2.3.4 CXCL7 (neutrophil-activating peptide-2)

CXCL7 is released by α -granules of platelets but is required to be activated by proteolytic cleavage by enzymes, for example, cathepsin-G (Walz, 1992). It also binds to CXCR2, therefore, it has chemotactic functions for both neutrophils and monocytes. There is an enhanced chemotactic activity of monocytes in response to CXCL7 in COPD patients (Traves et al., 2004). The

number of CXCL7⁺ cells, including endothelial cells, fibroblasts and submucosal glands in the bronchial submucosa are significantly higher in patients suffering from severe or very severe COPD compared to healthy non-smokers (Di Stefano et al., 2009a).

1.2.3.5 Other CXC related chemokines

Previous studies have shown that T cells, predominantly CD8⁺ T cells, play a potent role in the pathogenesis of COPD. Homing of T cells to the infected organs depends on initial activation followed by adhesion and selective chemotaxis. CXCL9 (monokine induced by interferon (IFN- γ), CXCL10 (IFN- γ -inducible protein of 10kDa, IP-10), and CXCL11 (IFN-inducible T-cell- α chemoattractant) are three chemokines released by IFN- γ activated bronchiolar epithelial cells and macrophages. They all can activate CXCR3 expressed by T cells and B cells (Devora, 2000). Studies have shown that CXCL10 expression is increased in bronchiolar epithelial cells and airway smooth muscle cells in COPD patients (Hardaker et al., 2004, Saetta et al., 2002), which leads to the accumulation of CXCR3⁺ CD8⁺ T cells (Tc1) and CD4⁺ T cells (Th1) in the epithelium and submucosa. CXCL11 has the highest affinity for CXCR3 (Clark-Lewis et al., 2003). Its

concentration is also increased in induced sputum of COPD patients.

CXCL12 (stromal-cell-derived factor-1) and CXCL13 (B-cell-attracting chemokine-1) are other two CXC chemokines binding to CXCR4 and CXCR5 respectively (Lapidot et al., 2005). CXCR4 is expressed by T cells, monocytes, immature B cells and airway epithelial cells upon activation. CXCR5 is predominantly expressed by mature B cells, T follicular helper cells (a subset of CD4⁺ helper T cells) and memory CD4⁺ T cells (Forster et al., 1996, Kim et al., 2001, Schaerli et al., 2000). Although their roles in COPD have not been explored yet, because their levels are increased in B cells and T cells associated with severe COPD, antagonists of CXCR4 and CXCR5 might be novel treatments for severe COPD.

CXCL16 (scavenger receptor for phosphatidylserine and oxidized low-density lipoprotein) is a transmembrane chemokine expressed by DCs, macrophages and B cells. It binds to CXCR6 expressed by Th1 cells (Kim et al., 2001a). Therefore, it could be relevant to COPD.

1.2.3.6 CCL2 (monocyte chemotactic protein-1, MCP-1)

CCL2 is a CC-chemokine and expressed by alveolar macrophages, T lymphocytes and epithelial cells. It is also released by type II pneumocytes in culture. Its secretion is triggered by LPS but inhibited by CSE. CCL2 activates CCR2 on monocytes and lymphocytes (Rose et al., 2003). Previous studies have shown that CCL2 levels are enhanced in induced sputum, BAL fluid and lungs of COPD patients (Capelli et al., 1999, de Boer et al., 2000, Traves et al., 2002). Recently, CCL2 expression has also been detected in small macrophages in induced sputum from COPD patients. Interestingly, there is a significantly increased level of mRNA expression of CCL2 in COPD patients compared to healthy non-smokers. Similar results are also seen in healthy smokers, although it is less pronounced (Frankenberger et al., 2010). TLR-2 and TLR-4 can down-regulate CCR2 expression (Parker et al., 2004). Blocking CCR2 induces the termination of the chemoattractant effect of induced sputum from COPD patients (Monzon et al., 2011), which indicates that antagonism of CCR2 may be a useful therapeutic strategy in COPD.

1.2.3.7 Other CC-chemokines

CCL3 (Macrophage inflammatory protein, MIP-1 α) and CCL4 (MIP-1 β) are secreted by macrophages, taken together with CCL5 (released by activated normal T cells expressed and

secreted, RANTES), they all not only activate CCR1 expressed by monocytes and neutrophils, but also bind to CCR5 expressed on Th1 and Tc1 cells (Devora, 2000). The levels of CCL3, CCL4 and CCL5 are all increased in COPD patients (Barnes, 2009). Numbers of CCR5⁺ T lymphocyte are elevated in induced sputum of COPD patients and in the airways of mild-moderate COPD patients but is decreased in severe disease (Costa et al., 2008, Di Stefano et al., 2001, Grumelli et al., 2004). CCL5 expression is increased in the airways and sputum of COPD patients during exacerbations, which indicates a role of CCR5 in the recruitment of T cells and macrophages during exacerbations (Fujimoto et al., 2005, Zhu et al., 2001a). In addition, CCL5 also activates CCR3 predominantly expressed by eosinophils.

CCL7 (MCP-3) and CCL13 (MCP-4) are CC-related chemokines and bind to CCR2 and CCR3 respectively. CCL17 (thymus- and activation-dependent chemokine, TARC), CCL20 (MIP-3 α) and CCL22 (macrophage-derived chemokine) are members of CC-chemokines and recruit target cells by interaction with CCR8, CCR6 and CCR4 respectively (Devora, 2000). Similar to CCL2, mRNA expression of CCL7, CCL13 and CCL22 are all markedly increased in small macrophages from induced sputum of COPD patients and healthy smokers compared to healthy non-

smoking subjects, although less pronounced in smokers (Frankenberger et al., 2010). CCL20 is a potent chemoattractant of dendritic cells and is mainly expressed by airway epithelial cells in response to IFN- γ (Reibman et al., 2003). Levels of CCL20 protein are increased in induced sputum from COPD patients compared to never smokers and smokers without COPD, and are negatively correlated with FEV₁% function. There is also a significant inverse correlation between CCL20 mRNA expression in total lung and FEV₁% (Demedts et al., 2007).

1.2.3.8 CX₃C chemokines

CX₃CL1 (fractalkine) is a unique CX₃C chemokine that is expressed on cell surfaces. Its expression is enhanced in human airway epithelial cells after IFN- γ induced stimulation. It is involved in the recruitment and adhesion of monocytes, T lymphocytes and natural killer cells to epithelial cells (Fujimoto et al., 2001). However, its role in COPD is unexplored. Recently it has been implicated in the function of CX₃CL1 in dendritic cell recruitment, which might be relevant to antigen presentation in COPD (Niess et al., 2005).

To sum up, cytokines and chemokine play potent roles in the pathogenesis of COPD. Figure 1.8 shows the pathological hallmarks of COPD.

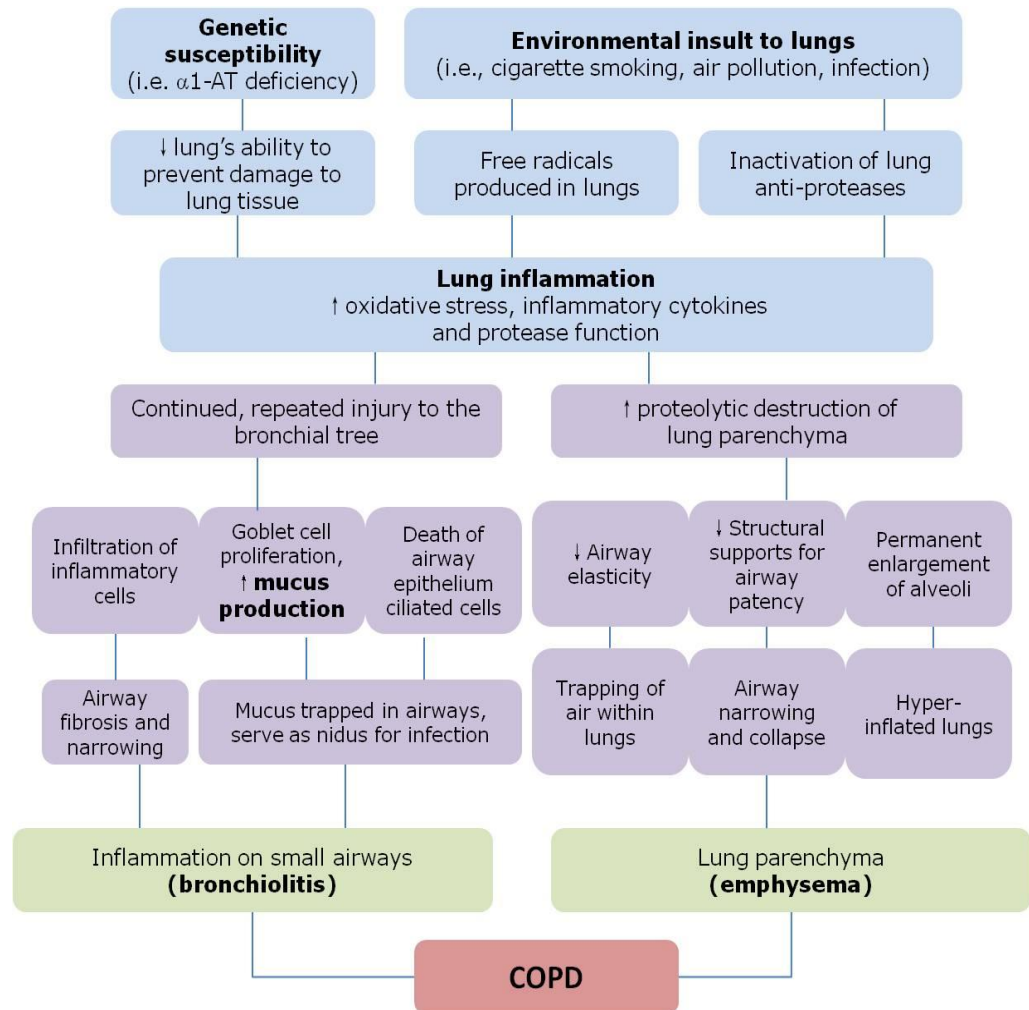


Figure 1.8 Pathological hall marks of COPD. The three main pathological hallmarks of COPD are chronic obstructive bronchiolitis, emphysema and mucus hypersecretion.

1.2.4 Oxidative stress in COPD

Excessively production of ROS can generate oxidative stress, which results in damage to lipids, proteins and DNA (Finkel, 2003). Oxidative stress can also lead to the oxidation of arachidonic acid and the formation of isoprostanes (a new series of prostanoid meidators), which have functional effects including bronchoconstriction and plasma exudation. Activated neutrophils, eosinophils, macrophages and epithelial cells are all good sources for ROS production in the airways (MacNee, 2001a). For examples, in neutrophils, hydrogen peroxide (H_2O_2) generated from superoxide anions ($\text{O}_2 \cdot^-$) is metabolized to hypochlorous acid in the presence of chloride ions, which is strong endogenous oxidant in the lung (Eiserich et al., 1998). Cigarette smoke, nitrogen dioxide, and ozone are exogenous oxidants that have the capacity to enhance the oxidative stress in the lungs (Pryor and Stone, 1993, Sunyer, 2001). In human airways, the production of oxidants is controlled by several antioxidant mechanisms (Cantin et al., 1990). However, the major antioxidants are extracellular, such as glutathione peroxidase that is significantly up-regulated in response to cigarette smoking and oxidative stress (Avissar et al., 1996).

There are abundant studies showing increased oxidative stress in COPD (MacNee, 2001b, Repine et al., 1997). Increased xanthine/xanthine oxidase activity has been observed in BAL fluid from COPD patients (Pinamonti et al., 1998). 4-hydroxy-2-nonenal is a specific marker for lipid peroxidation that is detectable in lungs of COPD patients and is location specific (Rahman et al., 2002). Ethane formed through lipid peroxidation is elevated in the breath of COPD patients and its concentration is correlated with disease severity (Paredi et al., 2000). Concentrations of H_2O_2 , 8-isoprostaglandin $\text{F}_{2\alpha}$, malondialdehyde and thiobarbituric acid reactive substances are all increased in exhaled breath condensate of COPD patients (Montuschi et al., 2000). Formation of ROS can activate NF- κ B and activator protein-1 transcription factors, which induce a neutrophilic inflammation through increased expression of IL-8 and other chemokines, TNF- α and MMP-9 (Rusznak et al., 1996). Therefore, this evidence indicates that increased oxidative stress in the respiratory tract plays a potent role in the disease by amplifying the inflammatory and the destructive responses in COPD.

1.2.5 Nitric oxide in COPD

Peroxynitrite is an oxidant and nitrating agent. It can damage a wide array of molecules in cells, including DNA and proteins

(Pacher et al., 2007). Increased oxidative stress leads to the formation of peroxynitrite via combination of nitric oxide (NO) with superoxide anions and then nitrate, which both change the gaseous phase of NO. This may explain why the NO levels are reduced in normal smokers (Kharitonov et al., 1995). However, in COPD patients, the NO levels are increased and are positively correlated with eosinophil numbers and bronchodilator response (Papi et al., 2000). The enzyme inducible NO synthase (iNOS) that generates NO is expressed in macrophages and lung parenchyma of patients with COPD, especially in severe COPD (Ichinose et al., 2000, Maestrelli et al., 2003). Furthermore, concentrations of nitrate are also elevated in breath condensate and sputum of smokers with or without COPD (Corradi et al., 2001, Kanazawa et al., 2003). Increased levels of NO are detected in the alveolar compartment of COPD patients and correlates with disease severity (Brindicci et al., 2005). This corresponds to the finding of increased inducible NO synthase in the periphery of lungs in COPD subjects (Maestrelli et al., 2003). Therefore, the peripheral NO may be a useful non-invasive biomarker of COPD inflammation (Barnes et al., 2006).

1.2.6 Peptide mediators in COPD

Endothelin-1 (ET-1) and bradykinin are two kinds of peptide related to COPD development (Barnes, 2004d). ET-1 is a potent vasoconstrictor with the ability to cause vascular smooth muscle hyperplasia (Bouallegue et al., 2007). Giaid *et al* have shown that expression of ET-1 is increased in pulmonary endothelial cells of COPD patients who have secondary pulmonary hypertension (Giaid et al., 1993). ET-1 concentrations are also increased in induced sputum and BAL fluid in COPD patients (Bacakoglu et al., 2003, Chalmers et al., 1999). All these findings indicate that ET-1 secretion may play an important role in pulmonary vasoconstriction and pulmonary hypertension in COPD. Bradykinin is a potent bronchoconstrictor of human airways, particularly small airways (Hulsmann et al., 1994). It can stimulate mucus secretion and potentiates cough in airways (Nagaki et al., 1996). Although its function in COPD is not fully explored, proinflammatory cytokines produced in COPD increase the expression of bradykinin receptors in pulmonary cells (Tsukagoshi et al., 1995).

1.2.7 Lipid mediators in COPD

1.2.7.1 Prostaglandin E₂ (PGE₂) and Prostaglandin F_{2α} (PGF_{2α})

PGE₂ is a bronchodilator of human airways (Pavord and Tattersfield, 1995). It inhibits the release of proinflammatory cytokines from monocytes and of acetylcholine from airway cholinergic nerves (Meja et al., 1997, Spicuzza et al., 1998). Therefore, PGE₂ may have beneficial effects in COPD. However, it can also stimulate mucus secretion and expression of mucin genes (Borchers et al., 1999), and activate airway sensory nerves to enhance coughing (Stone et al., 1992). PGF_{2α} is a bronchoconstrictor and has the ability of activating airway sensory nerves to produce cough (Nichol et al., 1990). Increased concentrations of both PGE₂ and PGF_{2α} are detected in exhaled breath of COPD patients (Montuschi et al., 2003).

1.2.7.2 Leukotriene B₄ (LTB₄)

LTB₄ is mainly released from activated alveolar macrophages and neutrophils in the airways (Shamsuddin et al., 1997). It is a potent chemoattractant of neutrophils and binds to its receptors, namely, BLT₁ and BLT₂, expressed on neutrophils and T lymphocytes respectively (Yokomizo et al., 2000). Increased levels of LTB₄ have been demonstrated in both exhaled breath condensate and induced sputum from patients with stable COPD (Beeh et al., 2003, Woolhouse et al., 2002), and particularly enhanced during exacerbations (Biernacki et al., 2003).

1.2.8 Growth Factors in COPD

The release of growth factors can induce fibrosis and cell proliferation and further cause structural changes in small airways and lung parenchyma.

1.2.8.1 Transforming growth factor- β (TGF- β)

TGF- β is a multifunctional growth factor with the ability to induce proliferation of fibroblasts and airway smooth muscle cells, deposition of extracellular matrix and epithelial repair. It also has immuno-suppressive effects on Th1, Th2 and Th17 cells (Yoshimura and Muto, 2011). Increased expression of TGF- β has been detected in airway epithelial cells and macrophages from small airways of COPD subjects (Takizawa et al., 2001). TGF- β levels are also elevated in peripheral blood monocytes from COPD patients (Hodge et al., 2003b). In peripheral lung tissue from COPD patients, TGF- β expression is increased and is correlated with immunoreactivity for 4-hydroxy-4-nonenal that is a marker of oxidative stress (Rahman et al., 2002).

1.2.8.2 Epidermal growth factor (EGF)

EGF, together with TNF- α , activates EGF receptor (EGFR) (Argast et al., 2004, Lee et al., 2007a). It plays a critical role

in regulation of mucus secretion, differentiation of mucus-secreting cells and expression of mucin genes (Burgel and Nadel, 2004). Increased levels of EGF, EGFR and the related cytokine, heregulin, have been detected in airway epithelial cells of COPD patients (Ganesan et al., 2012).

1.2.8.3 Vascular endothelial growth factor (VEGF)

VEGF is a regulator for the growth of new vessels and vascular leakage (Ho and Kuo, 2007). Its expression is increased in pulmonary vascular smooth muscle of patients with mild and moderate COPD but is reduced in severe COPD with emphysema (Santos et al., 2003). The level of VEGF is also reduced in induced sputum and the lung of COPD patients (Kanazawa and Yoshikawa, 2005).

1.2.8.4 Fibroblast growth factors (FGF)

FGF is related to vascular remodelling. Increased expression of FGF-1 and FGF-1 receptor, and FGF-2 have been investigated in large pulmonary vessels and small pulmonary vessels respectively (Kranenburg et al., 2005).

1.2.9 Proteases in COPD

1.2.9.1 Matrix metalloproteases (MMPs)

MMPs are a large family of up to 26 homologous endopeptidases, which can not only degrade protein components of the extracellular matrix but also has a potent function as regulating secretion of cytokines, chemokines and growth factors (Verma and Hansch, 2007). They are secreted by stromal cells and two major inflammatory cells including neutrophils and alveolar macrophages (Betsuyaku et al., 1999, Finlay et al., 1997a). Imbalance between proteinase and antiproteinase can lead to extracellular matrix destruction in the airways and contributing to airway remodelling and to decreased lung function in patients (Abboud and Vimalanathan, 2008). Therefore, the balance between MMPs and tissue inhibitors of MMPs (TIMPs) is thought to be crucial in disease pathogenesis (Barnes, 2002).

There are abundant studies showing the role of MMPs in COPD patients. MMP-9 is released by bronchial epithelial cells, neutrophils, alveolar macrophages and eosinophils (Atkinson and Senior, 2003). Its expression is induced by IL-13 and inhibited by TNF- α and IL-1 β . There are feedback loops between MMP-9 and TGF- β , and between MMP-9 and IL-8 (Finlay et al., 1998). The expression of MMP-9 on alveolar macrophages is elevated in the BAL fluid from patients with emphysema (Finlay et al., 1997b). Increased levels of MMP-9

and its inhibitor TIMP-1 are also detected in induced sputum from stable COPD patients, which correlates with both neutrophilia and FEV₁, compared to controls (Cataldo et al., 2000, Culpitt et al., 2005, Vernooy et al., 2004). In COPD exacerbations, levels of MMP-9 are markedly increased in induced sputum and are positively correlated with numbers of neutrophils and lymphocytes, but not macrophages (Mercer et al., 2005). Moreover, higher expression of MMP-9 is found in alveolar macrophages from healthy smokers than healthy non-smokers (Russell et al., 2002a).

A variety of experimental models of COPD have been established to detect effects of MMP-12 in COPD. Emphysema induced by cigarette exposure is prevented in MMP-12 knockout mice (Hautamaki et al., 1997). Also, MMP-12 ^{-/-} mice do not develop to the lavage neutrophilia as seen after smoke exposure. However, the neutrophil influx can be restored by instillation of wild type alveolar macrophages (Churg et al., 2003). Unlike findings in mice, studies on MMP-12 expression in human are controversial. Some studies demonstrate that MMP-12 mRNA is hardly detected in the lung (Gosselink et al., 2010, Imai et al., 2001, Lee et al., 2010), whereas others do show its difference between COPD patients and controls. For example, MMP-12 levels are higher in

induced sputum from stable COPD patients, no matter whether they are current smokers or ex-smokers, compared to healthy smokers and healthy non-smokers (Demedts et al., 2006, Ilumets et al., 2007). Its enzymatic activity is also significantly higher in COPD subjects. These data indicate that MMP-12 in sputum is not induced by cigarette smoking per se but by the disease itself. Babusyte *et al* show that the number of MMP-12 positive macrophages is higher in BAL fluid from current smokers with COPD than ex-smokers with COPD and controls (Babusyte et al., 2007). Another group also found the MMP-12 levels are higher in the macrophages from current smokers compared to ex-smokers (Wallace et al., 2008).

MMP-1 is also called interstitial collagenase-1. It is produced by bronchial epithelial cells, type II pneumocytes and alveolar macrophages in the lung (Verma and Hansch, 2007). Increased levels of MMP-1 mRNA and protein are detected in the lung of COPD patients (Gosselink et al., 2010, Lappalainen et al., 2005, Selman et al., 2003).

In addition, levels of MMP-2 are elevated in induced sputum, lung parenchyma and lung tissue of COPD patients (Gueders et al., 2006, Ramos et al., 2009, Segura-Valdez et al., 2000). Also, increased levels of MMP-8 are detected in induced

sputum, particularly during exacerbations in COPD patients (Ilumets et al., 2007, Skillrud et al., 1986, Vernooij et al., 2004).

Taken together, MMPs account for most of the elastolytic activity released from alveolar macrophages and neutrophils in COPD patients. They not only contribute to the destruction of extracellular matrix but also play important roles in small airway remodelling and emphysema.

1.2.9.2 Neutrophil elastase (NE)

NE is a serine protease that is inhibited by anti-protease α 1-antitrypsin in the lung parenchyma. It is stored in neutrophils and expressed on the cell surface when cells are activated by cytokines (Owen et al., 1997). NE is a potent mucus secretagogue of submucosal gland cells and goblet cells (Sommerhoff et al., 1990). It induces the expression of MUC5AC (Fischer and Voynow, 2002, Voynow et al., 1999). It also induces secretion of cytokines and chemokines, such as IL-8 (Nakamura et al., 1992). The apoptosis induced by macrophages are also impaired by NE (Vandivier et al., 2002). The amount of NE/ α 1-AT complexes is increased in BAL fluid

from current smokers with COPD and is negatively correlated with FEV₁ (Betsuyaku et al., 2000, Yoshioka et al., 1995).

1.3 Role of apoptosis in the pathogenesis of COPD

Unlike necrosis that is an uncontrolled process, apoptosis, programmed cell death, plays a key function in the elimination of unwanted damaged or infected cells (Baetu and Hiscott, 2002). Three major pathways have been involved in the regulation of apoptosis, namely, ligand-death receptor pathway (e.g, TNF- α and Fas ligand), cytolytic effector cell pathway (e.g, perforin and granzymes) and the mitochondrial intrinsic pathway (e.g. bcl-2). All pathways finally converge at the level of caspase-3 activation. Caspase activation leads to DNA fragmentation, characteristic of apoptotic cells and the cleavage of cytoskeletal proteins (Plataki et al., 2006). Figure 1.9 shows the different mechanisms of apoptosis induced by cytotoxic cells.

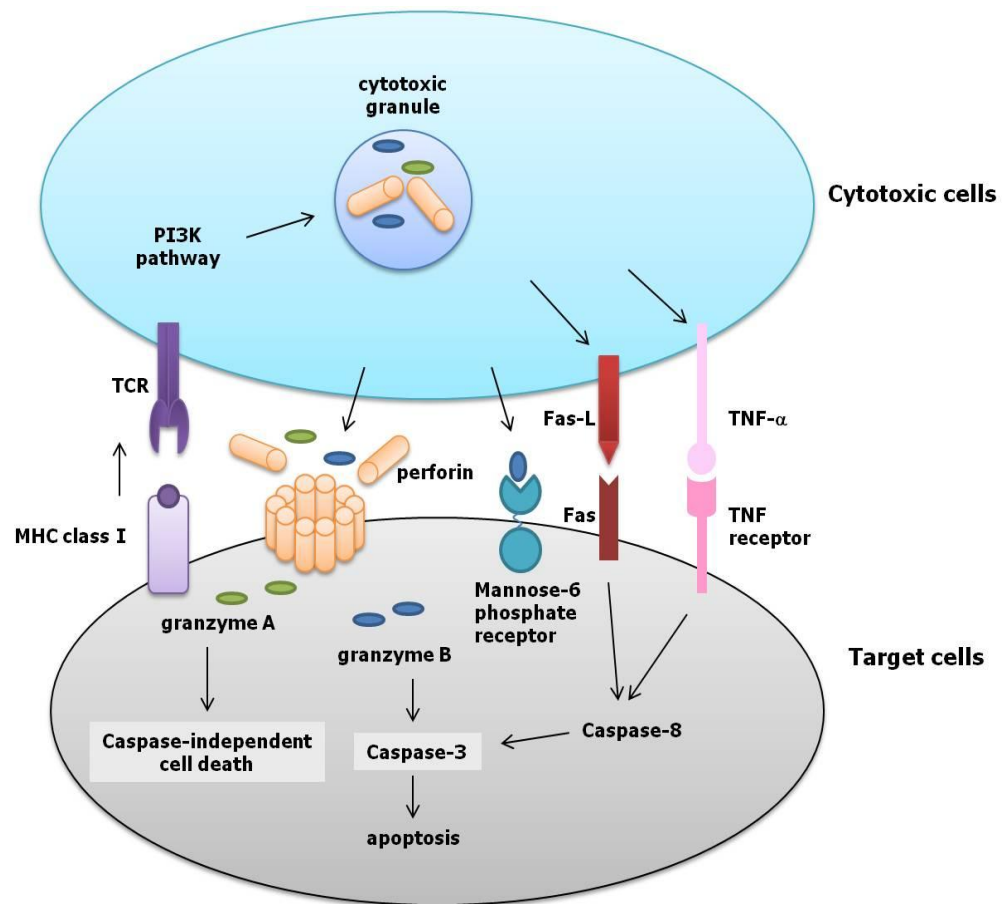


Figure 1.9 Different mechanisms of apoptosis induced by cytotoxic cells. Infected cells (Target cells) present antigen peptides to cytotoxic cells by binding to their TCRs. Activated cytotoxic cells then release granule killing molecules, such as perforin and granzymes. The former one is traditionally considered to form pores on the cytoplasm of target cells which facilitates granzymes passage. However, others indicate the granzyme B can still be endocytosised independently of perforin by mannose-6 phosphate receptors. Besides granule molecules, Fas-L and TNF- α expressed by cytotoxic cells interact with Fas and TNF receptor expressed by target cells respectively. Granzyme A and granzyme B induce cell apoptosis in different manners.

Studies on apoptosis in COPD are well established. Segura-Valdez et al were the first to show increased apoptosis in lung tissue sections of COPD patients compared to controls (Segura-Valdez et al., 2000). Both apoptosis and proliferation of cells (alveolar epithelial cells, endothelial cells and mesenchymal cells) have been elevated in the lung tissue of patients with emphysema compared to healthy subjects with or without smoking history (Imai et al., 2005, Yokohori et al., 2004). VEGF is a growth factor that induces the expression of anti-apoptotic proteins and is a survival factor of endothelial cells (Alon et al., 1995, Gerber et al., 1998). Its levels are decreased in induced sputum of COPD patients and correlated with disease severity. Moreover, both the percentage and absolute number of apoptotic bronchial epithelial cells were significantly higher in COPD patients, with increased expression of Fas, p53 and activated caspases (Hodge et al., 2005a). Reduced apoptosis of neutrophils was found in hospitalized patients with an acute exacerbation of COPD (Pletz et al., 2004).

In peripheral blood studies, several mediators involved in the induction of T cell apoptosis, such as TNF- α and Fas, are upregulated in COPD patients (Hodge et al., 2003b). There is a marked increase in soluble Fas in plasma from severe COPD

patients but not in mild or moderate COPD and healthy controls (Yasuda et al., 1998). Since COPD is a complex lung disease with systemic consequences, skeletal muscle apoptosis has been studied. Interestingly, the skeletal muscle apoptosis is elevated in COPD patients who have a low BMI (Agusti et al., 2002).

The major mechanism of cytotoxic cell-mediated killing is performed by cytotoxic granules and Fas-FasL (Russell and Ley, 2002). After recognition of target cells by effector cells, granules released by cytotoxic cells move to the immunological synapse by exocytosis. Then their membrane fuses with the killer cell plasma membrane and release contents into target cells by endocytosis (Hayes et al., 1989, Stinchcombe et al., 2001). A pore forming protein perforin and a group of serine proteases named as granzymes are two representative types of granules related to granule-mediated cytotoxicity (Pasternack et al., 1986, Podack et al., 1985).

1.3.1 Perforin

Perforin is homologous to the complement protein C9. It is expressed by all cytolytic cells including cytotoxic T cells, NK cells and NKT cells. Traditionally, perforin is thought to form pores in the target cell plasma membrane in a Ca^{2+} -dependent

manner that help granzymes to pass through (Tschopp et al., 1986). Indeed, pores are not big enough to allow passage of globular molecules as big as granzymes. Sublytic concentrations are required to deliver granzymes (Kawasaki et al., 2000, Metkar et al., 2002). Increased expression of perforin and granzyme B has been detected on CD8⁺ T cells from BAL fluid in COPD patients and healthy smokers compared non-smokers (Hodge et al., 2006). Percentages of killer cells expressing both perforin and granzyme B were also markedly higher in induced sputum of COPD groups (Chrysofakis et al., 2004, Urbanowicz et al., 2010). Morissette *et al* have shown that there is no significant difference in the percentage of CD8⁺ or CD4⁺ T lymphocytes and NK cells expressing perforin from peripheral blood between emphysematous smoking COPD patients and control groups (Morissette et al., 2007). Whereas Urbanowicz *et al* have demonstrated proportions of killer cells expressing both perforin and granzyme B are lower in the peripheral blood from COPD patients (Urbanowicz et al., 2009).

1.3.2 Granzymes

Granzymes are a family of highly homologous serine proteases. Their main function is to induce cell death to eliminate infected cells. Also, they play a role in immune regulation

(Regner and Mullbacher, 2004). Five human granzymes have been identified, namely, A, B, H, K, M (Barry and Bleackley, 2002). However, the most abundant granzymes are granzyme A and B, which induce cell death through alternate and nonoverlapping pathways.

Granzyme A

Granzyme A is the most abundant protease in the cytotoxic granules (Garcia-Sanz et al., 1990, Gershenveld and Weissman, 1986). It is mainly expressed by NK cells and activated cytotoxic T lymphocytes. Unlike perforin and other granzymes, granzyme A is continuously expressed after T cell activation, perhaps even in long-term memory cells (Chen et al., 2001, Zhang et al., 2003). Although it is also delivered into target cells via perforin, it initiates innate immune response via a granzyme B and caspase independent pathway and has the extracellular matrix cleavage capacities including degradation of type IV collagen (Chowdhury and Lieberman, 2008, Shresta et al., 1999). Interestingly, a recent study shows that, rather than its cytotoxic ability, the major role of granzyme A is promoting inflammation through inducing production of pro-inflammatory cytokines in monocytic cells (Metkar et al., 2008). Increased expression of granzyme A, but not granzyme B, has been detected in type II

pneumocytes in patients with severe COPD (Vernooy et al., 2007).

Granzyme B

Granzyme B is a serine protease and is classically considered to be expressed by cytotoxic T lymphocytes and NK cells (Fairclough et al., 2008). Recently, it has also been found in many different types of myeloid cells that are noncytotoxic, such as Treg cells, type II pneumocytes, basophils, macrophages, DCs, smooth muscle cells and so on, generally without perforin (Gondek et al., 2005, Grossman et al., 2004). Therefore, besides the major function in inducing cell death, granzymes also have additional non-cytolytic functions, for examples, regulating lymphocyte survival and immune tolerance, promoting inflammation and potentially enhancing lymphocyte migration (Chowdhury and Lieberman, 2008). Traditionally, perforin is thought to be essential in mediating the action of granzyme B against target cells. Perforin forms a multimeric transmembrane pore in the target cell, which facilitates granzyme B passing into the target cells (Bossi and Griffiths, 2005). Recently, another perforin-independent model of this process has been established. Granzyme B is endocytosed in complex with the mannose-6 phosphate receptor while perforin has not been expressed by effector

cells (Froelich et al., 1996, Metkar et al., 2002, Shi et al., 1997). Granzyme B mediated apoptosis is caspase-dependent. Caspase 3 is a central execution caspase and is directly processed by granzyme B (Talanian et al., 1997, Yang et al., 1998). In addition, activation of T cells induces the expression of perforin and granzymes. IL-2 is the only stimulus that consistently upregulates the expression of granzyme A, granzyme B and perforin (Liu et al., 1989). IL-15 and IL-21 also activate granzyme B and perforin expression in CD8⁺ T lymphocytes from human peripheral blood samples (White et al., 2007).

The specific role of granzyme B in COPD is much more studied than other granzymes. Hodge *et al* have shown increased expression of granzyme B and perforin by CD8⁺ T cells from BAL fluid in COPD patients and healthy smokers compared to non-smokers (Hodge et al., 2006), and there was a correlation between the percentage of granzyme B expressing T cells and bronchial epithelial cell apoptosis. In the same study, both soluble granzyme B and the percentage of T cells with intracellular granzyme B have been observed to be increased in COPD patients. Percentages of CD8⁺ T lymphocytes, NK cells and NKT-like cells expressing both perforin and granzyme B were markedly higher in induced sputum of COPD groups

(Urbanowicz et al., 2010). Nevertheless, findings produced using peripheral blood samples are complicated. Morissette *et al* have reported that there is no significant difference in the percentage of CD8⁺ or CD4⁺ T lymphocytes and NK cells expressing granzyme B between emphysematous smoking COPD patients and control groups including healthy non-smokers and healthy smokers (Morissette et al., 2007). Conversely, intracellular granzyme B has been investigated in a greater percentage of T cells from the peripheral blood of COPD patients (Hodge et al., 2006). Studies performed in our group have shown that the percentage CD8⁺ T lymphocytes, CD6^{dim}CD16⁺ NK cells and NKT-like cells that expressed both perforin and granzyme B were significantly lower in COPD patients compared with the healthy smokers and non-smokers (Urbanowicz et al., 2009).

Granzyme K

Granzyme K is another tryptase and causes cell death in a caspase-independent manner similar to granzyme A. However, it is expressed much less than granzyme A. Granzyme K is mainly expressed by activated killer cells and a small population of CD4⁺ T cells (Bratke et al., 2008, Hameed et al., 1988). Although there is no difference in granzyme K expression in BAL fluid from COPD patients and allergic

asthma (without allergen provocation), it is indeed upregulated in cases of acute airway inflammation (Bratke et al., 2008). These findings suggest that granzyme K release is a specific event in acute airway inflammation, and chronic inflammation, such as stable COPD, does not equate to a pan-granzyme effect and that there may be inter-granzyme variability regarding activity and levels in different diseases.

Other human granzymes

Up to date, there are no studies on the role of granzyme H and granzyme M in COPD.

1.3.3 Fas and Fas-ligand

Fas is a transmembrane receptor belonging to the TNF receptor family. It is widely expressed on normal cells and malignant cells (Leithauser et al., 1993, Nagata and Golstein, 1995, Smith et al., 1994). Fas ligand (Fas-L) is a membrane-bound cytokine belonging to the TNF family. It is mainly expressed on activated T cells and induces cell apoptosis by binding to Fas (Suda et al., 1995, Suda et al., 1993). Unlike levels of TNF- α that are significantly higher in peripheral blood from COPD patients, there is no significant difference in the serum level of soluble Fas-L and the plasma level of soluble Fas among groups (Takabatake et al., 2000). Interestingly,

another study has shown that the proportion of Fas positive T lymphocytes is dramatically higher in peripheral blood from COPD patients compared to healthy non-smokers and smokers. Moreover, the proportion of CD8⁺ T cells expressing Fas is correlated with disease severity (Domagala-Kulawik et al., 2007). Increased expression of Fas is also detected in bronchial epithelial cells in COPD subjects (Hodge et al., 2005a).

1.3.4 Granulysin

Granulysin is a proinflammatory molecule expressed by activated cytotoxic T cells and natural killer cells (Krensky and Clayberger, 2005). It is also a chemoattractant for monocytes, T lymphocytes, and activates the expression of many cytokines, such as IL-1, IL-6 and IL-10 (Deng et al., 2005). Studies have shown that granulysin has broad clinical relevance to a myriad disease including cancers (Kishi et al., 2002), skin afflictions (Ammar et al., 2008), infections (Sahiratmadja et al., 2007) and reproductive complications (King et al., 2003). However, it is still to be studied in COPD.

1.4 Hypothesis of this project

Chronic Obstructive Pulmonary Disease (COPD) is a chronic inflammatory disease involving both innate and adaptive immune responses. Activated inflammatory cells can release inflammatory mediators, such as $\text{TNF-}\alpha$, $\text{IFN-}\gamma$, matrix-metalloproteinase and IL-8, that sustain the inflammatory process and finally lead to airway remodelling (Chung, 2005), parenchymal destruction and mucociliary dysfunction (Danahay and Jackson, 2005). Activated neutrophils (Damiano et al., 1986, Galdston et al., 1977), macrophages (Molet et al., 2005, Montano et al., 2004) and DCs (Demedts et al., 2007) have been observed in the airways or sputum of COPD groups. CD8^+ T-lymphocytes, NKT-like cells and NK cells are the three main types of killer cells in the immune system and have been implicated in the pathogenesis of COPD (Fairclough et al., 2008). Abnormal numbers of CD8^+ T-lymphocytes have been found in large airways sub-epithelium (O'Shaughnessy et al., 1997, Lams et al., 2000), peripheral airway (Saetta et al., 1998), parenchyma (Saetta et al., 1999), smooth muscle (Baraldo et al., 2004), bronchial arteries (Peinado et al., 1999, Saetta et al., 1999, Turato et al., 2002), peripheral blood (Urbanowicz et al., 2009), sputum samples (Urbanowicz et al., 2010), and small airways such as bronchoalveolar lavage (BAL) (Saetta et al., 2002, Saetta et al., 2003). Abnormal proportions of NKT-like cells have also been detected in

peripheral blood (Urbanowicz et al., 2009), and induced sputum (Urbanowicz et al., 2010) from smokers with COPD. The frequency of iNKT cells is lower in peripheral blood from patients with exacerbations of COPD compared to stable COPD patients (Chi et al., 2011). Reduced numbers of NK cells and their phagocytic activities have been observed in peripheral blood from both asymptomatic smokers and COPD patients (Ferson et al., 1979, Prieto et al., 2001). However, NK cell proportion and their cytotoxicity are increased in induced sputum from COPD patients in contrast to healthy non-smokers and healthy smokers (Urbanowicz et al., 2010). Taken together, all these findings indicate that killer cells may be much more activated in COPD groups and play an important role in pathogenesis of disease. Figure 1.10 shows how NK cells interact with other leucocytes and epitheliums to become activated.

This project we are aiming to explore immunological features of killer cells in COPD, especially under the condition of cigarette smoking induced activation. Main approaches of this project are as followed. Firstly, we compared the activation status of killer cells *ex vivo* in both healthy participants and

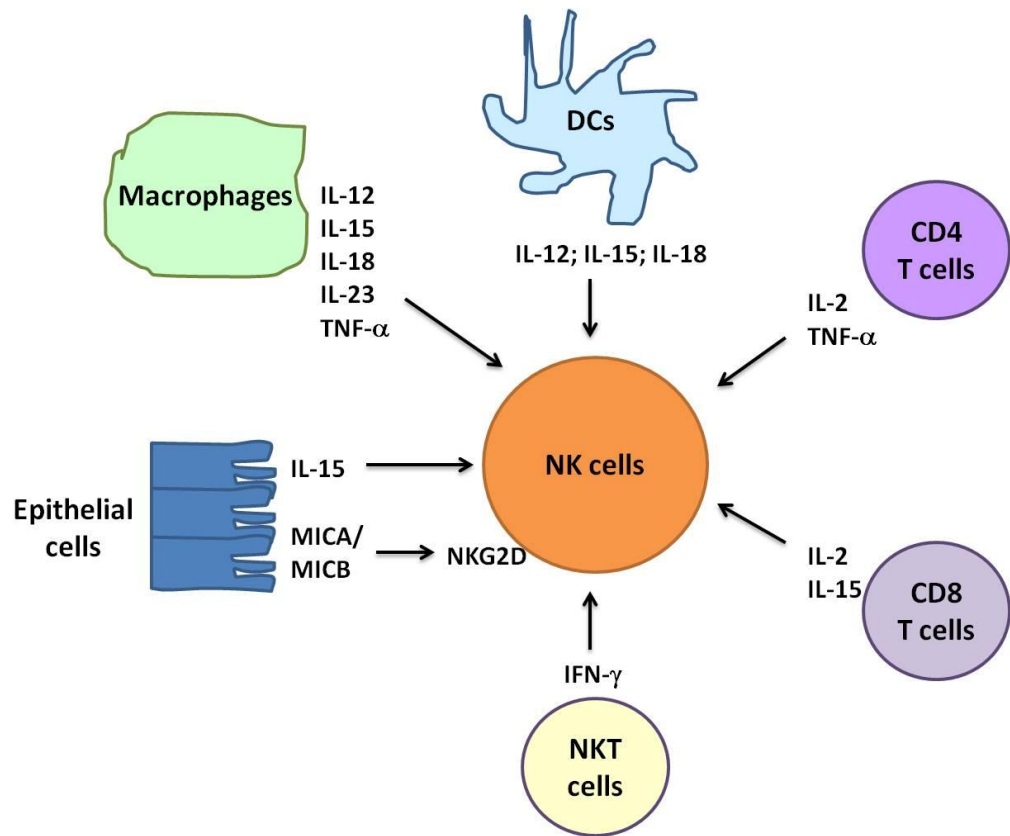


Figure 1.10 Interaction between NK cells and other lymphocytes. NK cells are the innate immune cells expressing multiple receptors. Activation of NK cells can be induced either by interaction of ligands to functional receptors, such as MICA to NKG2D; or by cytokines, for example, IL-2, 15, 18, 23, INF- γ and TNF- α , produced by other lymphocytes. Epithelial cell is a big source of producing IL-15.

COPD patients. Cigarette smoking is still the predominant risk factor in COPD, we hypothesis that cigarette smoke extract may induce immunological changes on killer cell functions. Therefore, *in vitro*, effects of cigarette smoke extract on killer cell activation upon different stimulation were investigated between study groups. Expression levels of effector molecules, such as granzymes, perforin and Fas-ligand, by killer cells

were also tested. Furthermore, cytotoxicity of CD8⁺ T cells in each study group was also investigated. All above research were performed in peripheral blood samples. Finally, we compared the activation levels of killer cells in BAL from COPD patients.

Material and Methods

2.1 Study population and procedures

The study population consisted of 16 healthy non-smoking volunteers, 12 healthy smokers, and 21 COPD patients who were either current smokers or ex-smokers. The diagnosis of COPD was made according to the ATS guidelines. Healthy non-smokers and current healthy smokers were with an $FEV_1 > 80\%$ of predicted normal and had normal lung function. COPD patients had irreversible airflow limitations on spirometry ($FEV_1 < 80\%$ of predicted with FEV_1/FVC ratio of $< 70\%$) and were reversibility to an inhaled beta-2 agonist of $< 10\%$ or < 200 mls absolute improvement. Smokers suffering from COPD accrued at least a 20-pack year smoking history and healthy smokers had a similar smoking history to them. All participants were age-matched and met the following inclusion criteria. Firstly, they were nonatopic (negative skin prick test response to common allergen extracts including grass pollen, house dust mite, cat dander and dog hair). Secondly, there was also no personal or family history of allergic diseases with them, such as allergic rhinitis. Thirdly, these subjects did not have any history of systemic or other pulmonary diseases, for example, TB, asthma, lung cancer, bronchiectasis or $\alpha 1$ -anti-trypsin deficient. In addition, they should also not be infected by acute viruses or bacteria. Finally, all COPD patients had been free of an acute exacerbation of COPD for at least 6

weeks preceding the study and none had received antibiotics or corticosteroids (oral and inhaled) over the same period.

Other four COPD patients were recruited for the BAL study. The inclusion/exclusion criteria were the same as mentioned above.

The study protocol was approved by the Nottingham Research Ethics Committee (REC reference 04/Q2403/102) and informed consent was obtained from all participants before entering the blinded study.

2.2 Peripheral Blood Mononuclear Cell (PBMCs) isolation

Peripheral venous blood (50mls) from each participant was diluted in warm RPMI 1640 (Sigma, Poole, UK) with 100U/ml penicillin and 100ug/ml streptomycin (RPMI-1640 medium) at 2:1 ratio and then layered over a discontinuous Histopaque density gradient (Sigma, Poole, UK). PBMCs were removed from the surface of Histopaque after centrifugation at 800g for 22 minutes and washed twice in RPMI medium at 300g, 4°C for 5 minutes. Isolated PBMCs were resuspended in appropriate medium. Prior to experiments, cells were counted with a Hemocytometer using standard protocol. Cell viability

was assessed with 0.5% Trypan Blue (Sigma, Poole, UK) staining.

2.3 CD56⁺ and CD8⁺ cell separation

Isolated PBMCs were incubated with anti-CD56 MicroBeads (Miltenyi Biotech Ltd, Surrey, UK) in MACS buffer containing sterile D-PBS (Sigma), 1% heat-inactivated fetal calf serum (FCS) (QB-Perbio, UK) and 0.4% EDTA (Sigma, Poole, UK) for 15 minutes at 4°C. CD56⁻ cell fractions were obtained by cell suspension passing through a cold MS column attached to a magnet. After washing three times, the column was removed from the magnet and 1ml MACS buffer was added and quickly plunged through the MS column into sterile tube for collection of CD56⁺ cells. To obtain NKT-like cells (CD3⁺CD56⁺) and NK cells (CD3⁻CD56⁺), CD56⁺ cells were incubated with Multisort Release Reagent for 10 minutes at 4°C, washed and then incubated with MultiSort Stop Reagent and α -CD3 microbeads for 15 minutes at 4°C. Finally the labelled cells were separated on a refrigerated LS column. CD8⁺ T lymphocytes were positively selected from CD56 negative fraction after incubated with anti-CD8 MicroBeads (Miltenyi Biotech Ltd, Surrey, UK) by using the same method of NKT-like cell separation. All isolated fractions were washed and counted. Detailed protocol for cell separation was shown on

manufacturer's instructions. The purity of CD3⁺CD56⁺, CD3⁻CD56⁺ cells or CD8⁺CD56⁻ cells was over 90%, as verified by flow cytometry.

2.4 Cigarette smoke extract (CSE) preparation

It should be noted that there are more than four thousand components in cigarettes that contribute to the diversity of cigarette brands. Choosing Marlboro to produce cigarette smoke extract in this project is because, of all menthol and non-menthol cigarettes, Marlboro is the brand used most often by 42.4% of smokers (Ref, office of applied studies. The NSDUH report Cigarette Brand Preferences in 2005. 2007).

A CSE collection apparatus is set up prior to 30 minutes of cell culture for each experiment in fume hood. As shown in Figure 2.1, the chemicals and equipment for the apparatus are as followed. (1) 10ml of Phenol-red free RPMI-1640 (GIBCO, Invitrogen, Paisley, UK) in a 30ml steralin; (2) cigarette (Marlboro, EU; Tar 10mg, Nicotine 0.8mg, carbon monoxide 10mg) with filter cut off; (3) 3-way tap; (4) two sections of flexible tubing; (5) steralin lid with hole cut in, for tubing to fit through; (6) 50ml syringe to aspirate the smoke; (7) two clamps and one clamp stand.

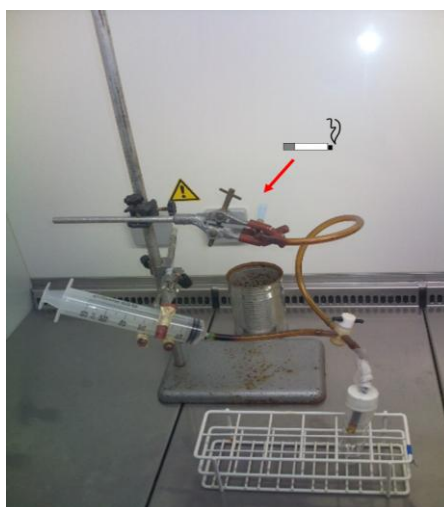


Figure 2.1 Apparatus of CSE preparation

To collect CSE, firstly, the tap should be switched to the position that keep the tubing leading to the sterilin is off. CSE from single lit commercial cigarette were aspirated to the 50ml syringe at rate of 125ml in 60s. Then, switch the tap so that the route to the sterilin containing 10ml of Phenol-red free RPMI-1640 is ON. Slowly bubble the entire 50ml of smoke into the medium. Repeat this process until the cigarette is finished. The crude CSE was filtered through a 0.45um pore filter to remove bacteria and/or large particles and subsequently adjusted with medium to an absorbance reading of 0.15 at 320nm. This solution was taken as 100% CSE.

2.5 Cell culture and treatments

Isolated PBMCs, NKT-like cells, NK cells or CD8⁺CD56⁻ T lymphocytes cells were resuspended in RPMI-1640 medium supplemented with 10% FCS at a density of 1×10^6 cells/ml. Cell suspension was loaded to a 96-well flat bottom tissue culture plate at 200ul of each well. To detect the cell activation, receptor expression and cytokine secretion, cells were stimulated with anti-CD3 (R&D System, Abingdon, UK), recombinant human IL-2 (rhIL-2; PeproTech, EC, London, UK), recombinant human IL-15 (rhIL-15; PeproTech, EC, London, UK), and CSE and incubated at 37°C in 5% CO₂. The time period of incubation and concentrations of these stimulators were determined by each experiment protocol. Specially, anti-CD3 was pre-coated in the plate for 90 minutes at 37°C in 5% CO₂ and wash in cold sterile D-PBS twice to remove the unbound antibody.

For cytotoxicity assay, isolated CD8⁺CD56⁻ cells were only incubated for 24 hours at 37°C in 5% CO₂. Before cell incubation, the plate was coated with 2µg/ml anti-CD3 (in 50ul D-PBS solution each well) for a minimum of 2 hours at 37°C in 5% CO₂. Target cells (P815 cells) were grown in RPMI-1640 medium supplemented with 10% FCS at 37°C in 5% CO₂ and ready for use when they reached confluence.

2.6 Extracellular staining and Intracellular staining

To harvest maximal cells for analysis, cells in each well were flushed with 200ul PBA (PBS; 1% BSA; and 0.1% sodium azide; Sigma, Poole, UK) twice and washed with 1ml PBA in a FACS tube (Elkay, Hampshire, UK). For extracellular staining, cells were incubated with fluorescently conjugated antibodies at the recommended concentrations for 30 minutes at 4°C. Unlabeled antibodies were removed with 1ml PBA washing up twice. Control tubes with single colour fluorescently conjugated antibody were used for setting compensation on flow cytometer by technician. Additional tubes with fluorescence minus one (FMO) were set up for quadrant setting in data presentation.

To detect the intracellular protein secretion, prior to 10 hours of harvesting, cells were treated with 10ug/ml brefeldin A (Sigma, Poole, UK). After washed with 1ml PBA solution, cell pellets were fixed in 3% formaldehyde in isotonic azide free solution, incubated for 5 minutes at 4°C, centrifuged, washed with 1ml PBA, 1ml PBA with 0.04% saponin (Sigma, Poole, UK) and 1ml PBA with 0.04% saponin and 10% FCS orderly. Interested labeled antibodies were mixed to cells at recommended concentrations and cells were incubated on a

rocker at 4°C for 2 hours (see table 1). Unbound antibodies were removed by 1ml PBA with 0.04% saponin washing up twice.

All centrifugations were performed at 300g for 5 minutes. Finally, cell pellets were fixed in 0.5% formaldehyde in isotonic azide free solution and stored at 4°C for flow cytometry analysis.

2.7 CD8⁺ T lymphocytes Redirected Cytotoxicity Assay

The cytotoxicity of CD8⁺ T lymphocyte was measured with colorimetric quantification of lactate dehydrogenase LDH released from lysed target cells (P815 cells). Harvested CD8⁺ CD56⁻ T cells were counted in a standard manner and resuspended at 4×10^6 cells/ml in RPMI-1640 medium without phenol red and FCS. P815 target cells were treated with 10ug/ml of anti-CD3 and incubated at 37°C. After 30 minutes incubation, P815 cells were washed twice and resuspended in RPMI-1640 without phenol red supplement with 10% FCS at a density of 1×10^5 cells/ml. A U-bottom 96-well culture plate was used in this protocol.

RPMI-1640 without phenol red supplement with 5% FCS was loaded to two triplicate sets of wells at 100ul/well. The top three wells, called as Culture Medium Background, were established to correct for contributions caused by phenol red and LDH activity that may be present in serum containing culture medium. The bottom three wells, called as Volume Correction Control, were used for correcting for the volume increase caused by addition of Lysis Solution (10X).

Target cells were added at 50ul/well to two triplicate sets of wells each containing 50ul RPMI-1640 without phenol red. The top three wells were used for measurement of target cell spontaneous LDH release, the bottom three was for the detection of target cell maximum LDH release.

Four effector to target ($E:T$) ratios were tested in this assay, namely, 40, 20, 10 and 5. Diluted effector cells at 50ul were loaded to each well containing 50ul of RPMI-1640 without phenol red supplement with 10% FCS. Target cells at 50ul were mixed with 50ul of diluted CD8⁺ T cells. Both conditions were prepared in triplicates as well.

Effector and target cells were brought into close contact by centrifuging the plate at 250g for 4 minutes and incubated for

4 hours at 37°C in 5% CO₂. Forty-five minutes prior to harvesting supernatants, wells used for Volume Correction Control and target cell maximum LDH release were treated with Lysis Solution (10X) at 10ul/well. After 4 hours incubation, plate was centrifuged at 250g for 4 minutes. 50ul aliquots from all wells were transferred to a fresh 96-well flat-bottom (enzymatic assay) plate. Assay buffer Substrate Mix (Promega, USA) was thawed at 37°C shortly and added to each well at 50ul to detect LDH activities. The plate was then covered with foil and incubated for 30 minutes at room temperature. After incubation, 50ul of Stop Solution (Promega, USA) was added to each well and the sample was read at 49nm on an Emax precision microplate reader (Molecular Devices, Workingham, UK) using SOFTmax software (Molecular Devices) immediately. The percentage of cell-mediated cytotoxicity was calculated by the following formula:

$$\text{Cytotoxicity (\%)} = \left\{ \frac{(\text{effector/target cell mix-spontaneous effector LDH release} - \text{spontaneous target LDH release})}{(\text{maximum target LDH release} - \text{spontaneous target LDH release})} \right\} \times 100$$

2.8 BAL preparation

Bronchoscopy with BAL was performed by a clinician from the Queen's Medical Centre, University of Nottingham, UK. Four COPD patients were recruited to this research. Each of them was sedated with oxycodon and scopolamine hydrobromide 10 minutes prior to the procedure. A fibreoptic bronchoscope (Olympus 1T 20 D, Olympus Corporation, NY, USA) was pushed in the subsegmental bronchus of the right middle lobe via the nasal route. Four 50ml aliquots of sterile, pre-warmed normal saline were administered and sucked into four plastic bottles by gentle hand suction through the bronchoscope. All bottles were kept in ice. The first pooled samples were discarded. Other three samples were used for BAL cellular analysis.

Before centrifugation of BAL fluid, two 50ml centrifuge tubes were weighted on an electronic scale. Samples were pipetted into the tubes and the volumes of them were noted. Filled tubes were re-weighted and the weight of recovered BAL fluid was calculated. A sterile piece of the 48um nylon gauze was put in a 35mm sterile funnel to pour through BAL samples, and then the volume and the weight of filtrate were measured and noted. Filtrated BAL fluid was centrifuged at 400g for 10 minutes at 4°C. The cell pellets were washed twice with sterile D-PBS by centrifugation. Total cell counts were determined

with a haemocytometer. Cells were resuspended in RPMI medium supplemented with 10% FCS. To determine the cell populations in the BAL, extracellular staining was performed for these cells as described previously.

2.9 Flow cytometric analysis

Labelled cells were analyzed on an EPICS ALTRA flow cytometer (Beckman Coulter, Buckinghamshire, UK). For each test, 50000 total events were collected, unless otherwise stated. Data analysis was performed using WEASEL software version 2.3 (The Walter And Eliza Hall Institute of Medical Research, Melbourne, Australia). For each staining experiments, compensation detection was set up by using control tubes with single colour fluorescently conjugated antibody. Fluorescence minus one (FMO) tubes were also used for quadrant setting. Dead cells were excluded according to forward scatter vs side scatter characteristics.

2.10 Statistical analysis

The statistical analysis was performed with Prism software, version 4.0c (GraphPad). Data that were non-normally distributed are expressed as median and range, unless otherwise stated. Normality was detected using the Kolmogorov–Smirnov test. Differences between the four

groups of participants were tested using the non-parametric Kruskal-Wallis test with *post hoc* pairwise comparisons made by the Dunn's Multiple Comparison test to determine which pair was statistically significantly different. Differences between the "before and after" treatment were tested by parametric student's *t* test. Confidence interval 99% was used to show two-tailed *p* values. *P* values of less than 0.05 or 0.01 were considered to indicate statistical significance according to the size of each study group.

2.11 Antibodies for flow cytometry

Table 2.1 shows all antibodies used in whole project. FITC, PE and PC7 labelled antibodies were added at 5ul to cell pellets; ECD and PC5 were added at 2.5ul.

Table 2.1 Antibodies for extracellular staining and intracellular staining

Antigen	Fluorochrome	Isotype	Clone	Source
CD3	ECD PC7	Mouse IgG1	UCHT1	Beckman Coulter, Luton, UK
CD4	FICT PC5	Mouse IgG1	13B8.2	Beckman Coulter, Luton, UK
CD8	PC5	Mouse IgG1	B9.11	Beckman Coulter, Luton, UK
CD8	ECD	Mouse IgG1	SFC121Thy2D3	Beckman Coulter, Luton, UK
CD25	FITC PE	Mouse IgG2a, k	B1.49.9	Beckman Coulter, Luton, UK
CD56	PE PC5	Mouse IgG1	N901	Beckman Coulter, Luton, UK
CD69	ECD PC5	Mouse IgG2b	TP1.55.3	Beckman Coulter, Luton, UK
KIR3DL1 (CD158e1)	FITC	Mouse IgG1, k	DX9	Biolegend, San Diego, CA, USA
NKG2D (CD314)	PE	Mouse IgG1	ON72	Beckman Coulter, Luton, UK
GranzymeB	FITC	Mouse IgG1, k	GB11	BD Pharmingen™
Perforin	PE	Mouse IgG2b, k	δG9	BD Pharmingen™
Fas-L (CD178)	PE	IgG Hamster	4A5	Beckman Coulter, Luton, UK

Results

3.1 Activation of killer cells in peripheral blood in COPD patients *ex vivo*

3.1.1 Introduction

Many risk factors have been implicated in COPD, but cigarette smoking is still considered the common and key pathological driver in the development and progression of COPD (Mannino and Buist, 2007). Smoking cessation has shown great potential in control of disease development (Scanlon et al., 2000, Tashkin and Murray, 2009).

CD8⁺ T-lymphocytes, NKT-like cells and NK cells are the three main types of killer cells in the immune system and have been implicated in the pathogenesis of COPD (Fairclough et al., 2008). Abnormal numbers of CD8⁺ T-lymphocytes have been found in large airways sub-epithelium (O'Shaughnessy et al., 1997, Lams et al., 2000), peripheral airway (Saetta et al., 1998), parenchyma (Saetta et al., 1999), smooth muscle (Baraldo et al., 2004), bronchial arteries (Peinado et al., 1999, Saetta et al., 1999, Turato et al., 2002), peripheral blood (Urbanowicz et al., 2009), sputum samples (Urbanowicz et al., 2010), and small airways such as bronchoalveolar lavage (BAL) (Saetta et al., 2002, Saetta et al., 2003). Abnormal proportions of NKT-like cells have also been detected in

peripheral blood (Urbanowicz et al., 2009), and induced sputum (Urbanowicz et al., 2010) from smokers with COPD. The frequency of iNKT cells is lower in peripheral blood from patients with exacerbations of COPD compared to stable COPD patients (Chi et al., 2011). Reduced numbers of NK cells and their phagocytic activities have been observed in peripheral blood from both asymptomatic smokers and COPD patients (Ferson et al., 1979, Prieto et al., 2001). However, NK cell proportion and their cytotoxicity are increased in induced sputum from COPD patients in contrast to healthy non-smokers and healthy smokers (Urbanowicz et al., 2010). In addition, cigarette smoke treated NK cells from non-smoking healthy subjects have shown defective cytokine secretion upon poly I:C activation (Mian et al., 2008).

Studies on CD4⁺ T cells are less extensive than CD8⁺ T cell infiltrates. CD4⁺ T cells have been shown to be increased in airways and lung parenchyma in COPD patients (O'Shaughnessy et al., 1997, Saetta et al., 1999), particularly in proximity to bronchus associated lymphoid tissue (Hogg et al., 2004). CD4⁺ helper T cells are mainly responsible for orchestrating downstream immune process by producing cytokines. They are essential for the full development of

adaptive immune cytotoxicity by priming and promoting the long-term survival of CD8⁺ T cells.

Activation is a prerequisite for killer cells to be functional. Activated killer cells can cause the production of inflammatory cytokines, chemokines and cytotoxic molecules, such as INF- γ , TNF- α , IL-8 and granzymes, which can induce pathological features including emphysema, mucus hypersecretion and cell apoptosis, and lung tissue damage in COPD (Barnes, 2004d). Several studies have investigated expression of activation markers on T-lymphocytes from peripheral blood, induced sputum and bronchial alveolar lavage (BAL) fluid from healthy smokers and COPD patients with varying results (Glader et al., 2006b, Leckie et al., 2003), but none have looked specifically at CD8⁺ T lymphocytes, NKT-like and NK cells. CD69 and CD25 are cell surface markers of early and late stages of lymphocyte activation, respectively (Caruso et al., 1997). Killer cell activation involves several factors, such as interaction between antigen receptors and peptides presented by MHC class I molecules or MHC class I-related proteins. Human killer cells express immunoglobulin-like receptors (KIRs), belonging to the Ig superfamily (CD158 family), which bind to MHC Class I. They can be classified into two groups, namely, inhibitory KIRs and activating KIRs. Inhibitory KIRs

possess long (L) cytoplasmic domains, such as KIR2DL or KIR3DL. Activating KIRs possess short cytoplasmic domains with an immunoreceptor tyrosine-based activation motif (ITAM)-bearing DAP12 adapter protein (Parham, 2005). NKG2D is a unique activating receptor and a type II transmembrane-anchored glycoprotein. It transmits signals through its transmembrane segment associating with adaptor protein DAP10. In the absence of DAP10, NKG2D is not able to be presented on cell surface and is retained in the cytoplasm (Diefenbach et al., 2002, Wu et al., 1999). Expression of NKG2D can be regulated by cytokines, for example, IL-15 and TNF- α can enhance NKG2D expression whereas TGF- β causes its downregulation (Castriconi et al., 2003, Lee et al., 2004, Parasa et al., 2012). In humans, KIR3DL1 (CD158e1) and NKG2D are two representative signaling receptors expressed by CD8⁺ T-lymphocytes, NK cells, NKT cells and $\gamma\delta$ TCR⁺ T cells (Ferrini et al., 1994, Moretta et al., 1990, Patterson et al., 2008, Phillips et al., 1995, Wu et al., 1999). They can recognize MHC class I molecules and MHC class I-related proteins, respectively.

In this chapter, the activation levels (by measurement of CD69 and CD25 expression) of the three main classes of human killer cells, namely, CD8⁺ T-lymphocytes, NKT-like

cells and NK cells in peripheral blood were investigated. Samples were obtained from four groups of participants including healthy non-smoking volunteers, current healthy smokers, current smokers with COPD and ex-smokers with COPD. To further detect the association between cell activation and their functional receptors in peripheral blood samples, we also examined the expression of the inhibitory receptor KIR3DL1 (CD158e1) and the activating receptor NKG2D in these four groups.

3.1.2 Material and methods

3.1.2.1 Study population and procedures

The Nottingham Research Ethics Committee approved the study protocol (REC reference 04/Q2403/102) and written informed consent was obtained from all participants before entering the study. Thirty-two participants were included in both the peripheral blood activation and the KIR study. Participants were diagnosed as having COPD according to the ATS guidelines, were either current smokers with COPD (cuS-COPD) or ex-smokers with COPD (exS-COPD) and had accrued at least a 20-pack year smoking history. Healthy non-smokers (HNS) and current healthy smokers (HS), with an FEV₁ above 80% of predicted, were matched, as far as possible, for age, with participants with COPD; healthy smokers were also

matched for smoking history. Table 3.1.1 details the demographic and spirometric data of the participants in this study. Participants were excluded as previously stated ; specifically, they were all non-atopic (negative skin prick test response to common allergen extracts including grass pollen, house dust mite, cat dander and dog hair) and all COPD patients had been free of an acute exacerbation of COPD for at least 6 weeks preceding the study. In addition, none had received antibiotics or corticosteroids (oral and inhaled) over the same period.

3.1.2.2 Peripheral Blood Mononuclear Cell isolation

PBMCs were isolated from whole blood on a discontinuous histopaque density gradient (Sigma, Poole, UK). Briefly, 50ml of whole blood were diluted 2:1 in RPMI 1640 medium (Sigma, Poole, UK), were diluted 2:1 in RPMI 1640 medium (Sigma, Poole, UK), layered over histopaque and centrifuged for 22 minutes at 800g. The resultant mononuclear layer was removed and washed twice and resuspended. Cells were counted with a Hemocytometer using trypan blue.

Table 3.1.1 Demographic and spirometric values of the studied groups

	Healthy non-smokers	Current healthy smokers	cuS-COPD	exS-COPD
Participants	8	8	9	7
Age (years)	54 (47-66)	53 (45-63)	61 (45-77)	62 (49-69)
Gender (M/F)	4/4	3/5	3/6	3/4
Packs/yrs	0	35 (32-40)	56 (30-77)	36 (20-51)
Smoking status (Current/Ex)	0	8/0	9/0	0/7
Chronic bronchitis (Yes/No)	0	3/5	6/3	5/2
FEV₁ (% pred)	105 (75-126)	101 (85-114)	41 (30-61)	58 (52-66)
FEV₁/FVC (%)	104 (67-120)	89 (66-101)	48 (29-53)	51 (36-65)
ΔFEV₁ post bronch	2.2 (1.9-3.0)	2.1 (1.7-2.9)	1.6 (1.1-2.4)	1.1 (0.6-1.5)
BMI (kg/m²)	22.8 (18.7-26.2)	24.7 (19.2-31.6)	23.2 (18.3-30.8)	21.5 (19.6-23.1)
Inhaled GCS (on/off)	N/A	N/A	3/6	4/3
MRC dyspnoea scale	N/A	N/A	3 (2-4)	2 (1-4)
Distance walked in 6 min (m)	N/A	N/A	308 (50-431)	277 (140-456)
BODE Index	N/A	N/A	5 (2-7)	5 (1-6)

Results are expressed as median with range in brackets.

exS-COPD, ex-smokers with COPD; cuS-COPD, current smokers with COPD; FEV₁, forced expiratory volume in 1 second; pre, predicted value; FVC, forced vital capacity; BMI, body mass index; GCS, corticosteroid; MRC, medical research council; BODE, **B**MI, **O**airflow **O**bstruction, **D**yspnoea and **E**xercise capacity.

3.1.2.3 Flow cytometric analysis

Cells were stained for flow cytometry as previously described . Briefly, cells were fixed in 3% formaldehyde in isotonic azide free solution (Beckman Coulter, Luton, UK) and washed before labelled antibodies (Table 3.1.2) were added at the recommended concentration. The cells were incubated in the dark and excess antibody removed. Flow cytometric analysis of antibody labelled cells was performed using an EPICS Altra (Beckman Coulter, Luton, UK). Fifty thousand live-gated events were collected for each sample and isotype matched antibodies were used to determine binding specificity. Data were analysed using WEASEL version 2.3 (WEHI). Necrotic cells were excluded from analysis according to their forward and side scatter characteristics.

3.1.2.4 Statistical analysis

The statistical analysis was performed with Prism software, version 4.0c (GraphPad). Normality was detected using the Kolmogorov–Smirnov test. All data are expressed as median and range. Differences between the four groups of participants were tested using the non-parametric Kruskal-Wallis test with *post hoc* pairwise comparisons made by the Dunn's Multiple Comparison test to determine which pair was

statistically significantly different. P values of less than 0.05 were considered to indicate statistical significance.

Table 3.1.2 Antibodies used for flow cytometry

Antigen	Fluorochrome	Isotype	Clone	Source
CD3	ECD PC7	Mouse IgG1	UCHT1	Beckman Coulter, Luton, UK
CD8	PC5 APC	Mouse IgG1	B9.11	Beckman Coulter, Luton, UK
CD8	ECD	Mouse IgG1	SFC121Thy2 D3	Beckman Coulter, Luton, UK
CD25	FITC PE	Mouse IgG2a	B1.49.9	Beckman Coulter, Luton, UK
CD56	PE PC5 PC7	Mouse IgG1	N901	Beckman Coulter, Luton, UK
CD69	ECD PC5	Mouse IgG2b	TP1.55.3	Beckman Coulter, Luton, UK
KIR3DL1 (CD158e1)	FITC	Mouse IgG1	DX9	Biolegend, San Diego, CA, USA
NKG2D (CD314)	PE	Mouse IgG1	ON72	Beckman Coulter, Luton, UK

3.1.3 Results

3.1.3.1 Demographic and medical characteristics of COPD participants, current healthy smokers and healthy controls

There were no statistical difference between groups in terms of age (Figure 3.1.1.A) or pack years smoked (Figure 3.1.1.B) between ex-smokers with COPD, current smokers with COPD and current healthy smokers. Furthermore, there was no statistical difference in the inhaled corticosteroid use between ex-smokers with COPD and current smokers with COPD.

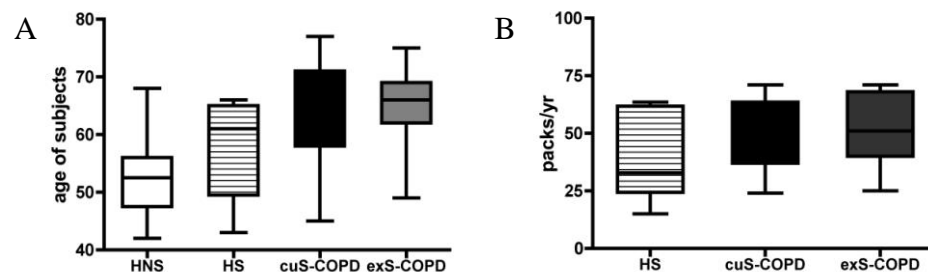


Figure 3.1.1 Comparison of ages and pack year smoked between interested groups. There were statistical differences in terms of (A) ages or (B) pack year smoked between healthy non-smokers (HNS, n=8), current healthy smokers (HS, n=9), current smokers with COPD (cuS-COPD, n=9) and ex-smokers with COPD (exS-COPD, n=7).

3.1.3.2 Killer cell activation in peripheral blood

All individuals had similar total mononuclear cell numbers, which were within the normal range (data not shown). Representative flow analysis dot plots of Forward Scatter (FS) vs Side Scatter (SS) for identification of live and dead cells (Figure 3.1.2.A), along with dot plots for identification of CD8⁺ T-lymphocytes (CD3⁺CD8⁺), CD4⁺ T-lymphocytes (CD3⁺CD4⁺) (Figure 3.1.2.B), NKT-like cells (CD3⁺CD56⁺), and NK cells (CD3⁻CD56⁺) (Figure 3.1.2C) are shown. In addition, exemplar dot plots for CD69 and CD25 staining of these gated four cell types from peripheral blood in the four groups are shown in Figure 3.1.3. Cells are designated activated when they express one or both of the activation markers (i.e., CD69 positive, CD69/CD25 double positive or CD25 positive).

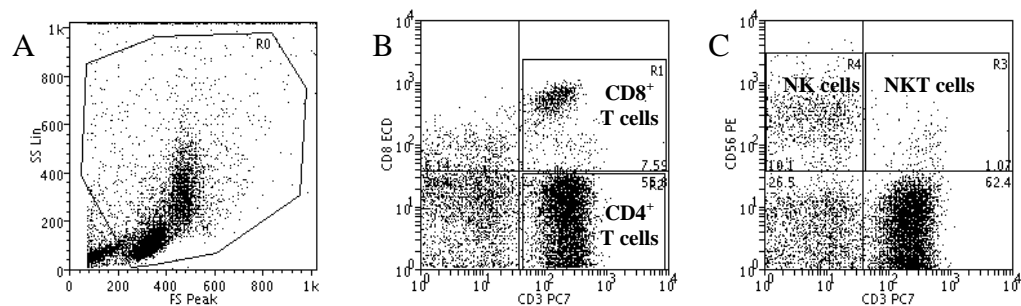
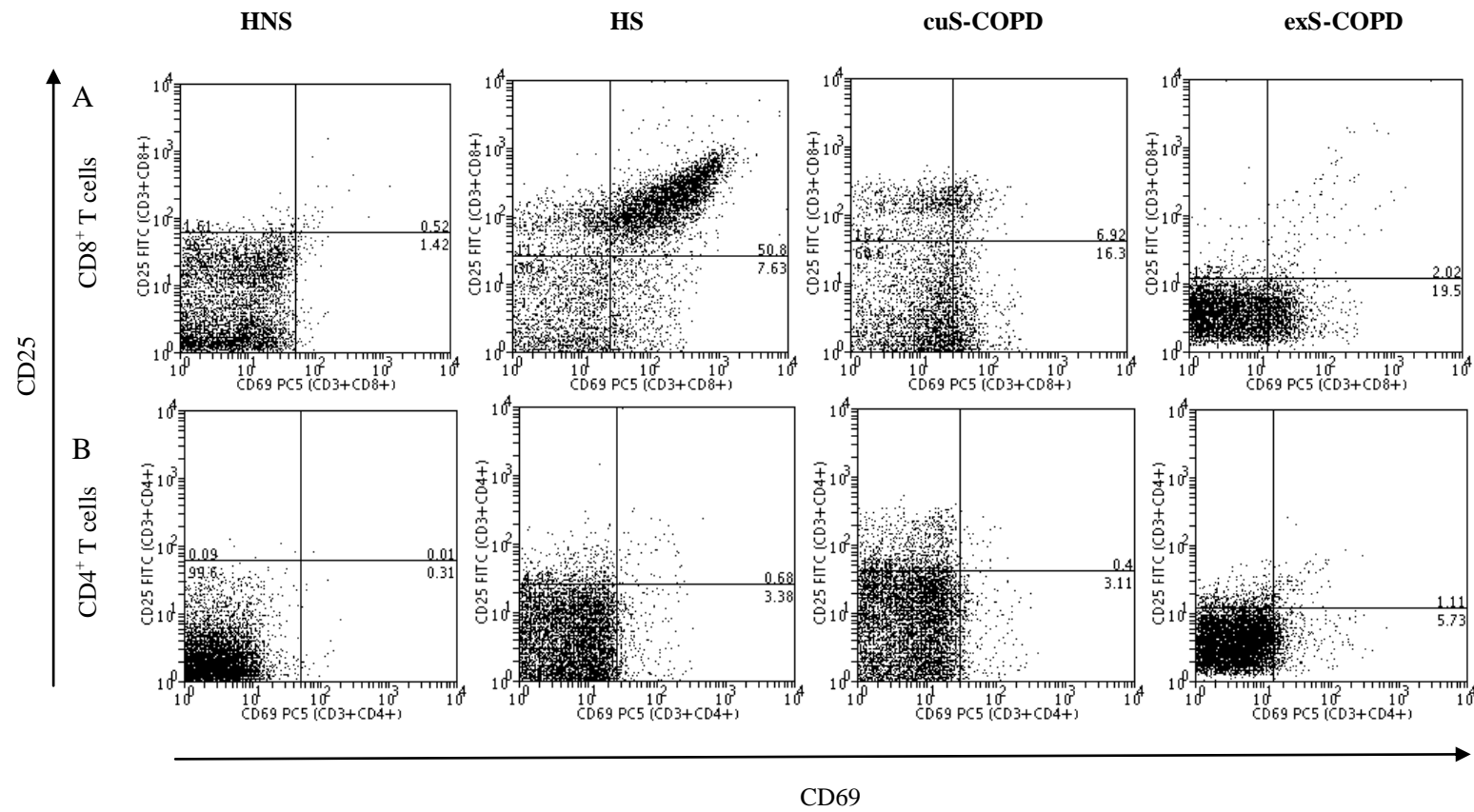


Figure 3.1.2 Representative dot plots gating for live/dead cells, identification of cell populations from current smokers with COPD. (A) Forward Scatter (FS) vs Side Scatter (SS) plot for identification of live and dead cells; (B) and (C), Identification of CD8⁺ T cells (CD3⁺CD8⁺), CD4⁺ T cells (CD3⁺CD4⁺), NKT-like cells (CD3⁺CD56⁺) and NK cells (CD3⁻CD56⁺).



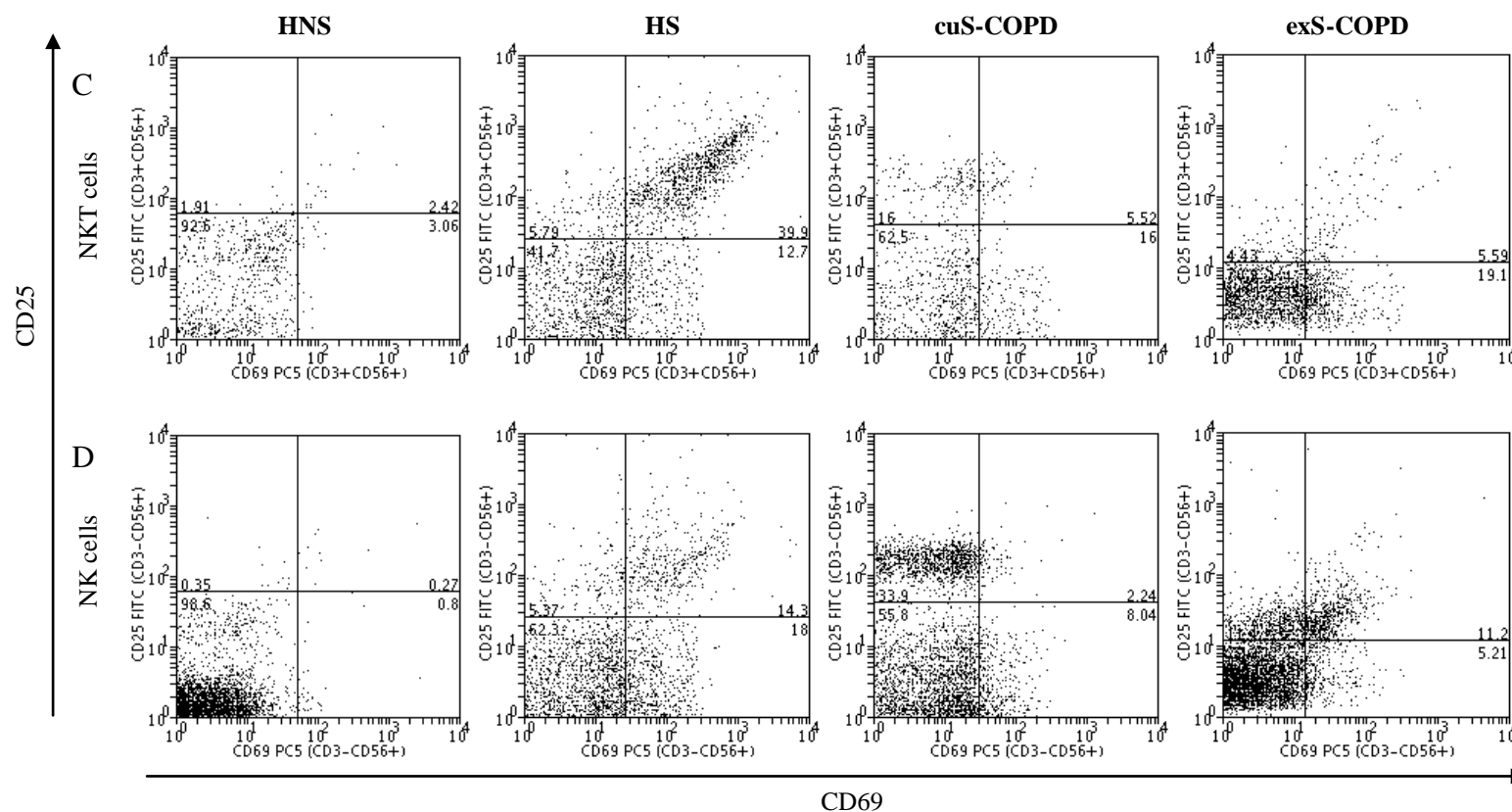


Figure 3.1.3 Representative dot plots for activation of interested cell types from four groups including healthy non-smokers (HNS), current healthy smokers (HS), current smokers with COPD (cuS-COPD) and ex-smokers with COPD (exS-COPD). Activation is designated by expressing either CD69 alone (lower right quadrant), CD69 and CD25 (upper right quadrant) or CD25 alone (upper left quadrant). (A) CD8⁺ T cells; (B) CD4⁺ T cells; (C) NKT-like cells; and (D) NK cells.

Significantly more CD8⁺ T-lymphocytes from current healthy smokers (41.96%, 27.2-72.8; $p<0.001$), current smokers with COPD (34.3%, 13.8-56.2; $p<0.01$), and ex-smokers with COPD (23%, 11.01-48.4; $p<0.05$) were activated *ex vivo* compared to healthy non-smokers (3.78%, 2.5-9.3) (Figure 3.1.4.A). Analyses of individual markers (i.e., CD69 positive/CD25 negative; CD69 negative/CD25 positive; or CD69 positive/CD25 positive) were not different between groups or cell types (data not shown).

NKT-like cells showed a similar pattern to CD8⁺ T-lymphocytes, namely significantly more *ex vivo* activation from both current healthy smokers (40.05%, 17.7-58.7; $p<0.001$) and current smokers with COPD (33.04%, 15.7-48.8; $p<0.01$) compared to healthy non-smokers (7.3%, 4.26-16.82) (Figure 3.1.4.C). Also, no significant differences in NKT-like cell activation were observed between ex-smokers (27.05%, 12.4-48.1) and healthy non-smokers (7.3%, 4.26-16.82; $p>0.05$), current healthy smokers (40.05%, 17.7-58.7; $p>0.05$), or current smokers with COPD (33.04%, 15.7-48.8; $p>0.05$).

NK (CD56⁺CD3⁻) cells from both current healthy smokers (39.55%, 14.78-50.1; $p<0.05$) and current smokers with COPD (44.18%, 14.2-72.8; $p<0.01$) were significantly more

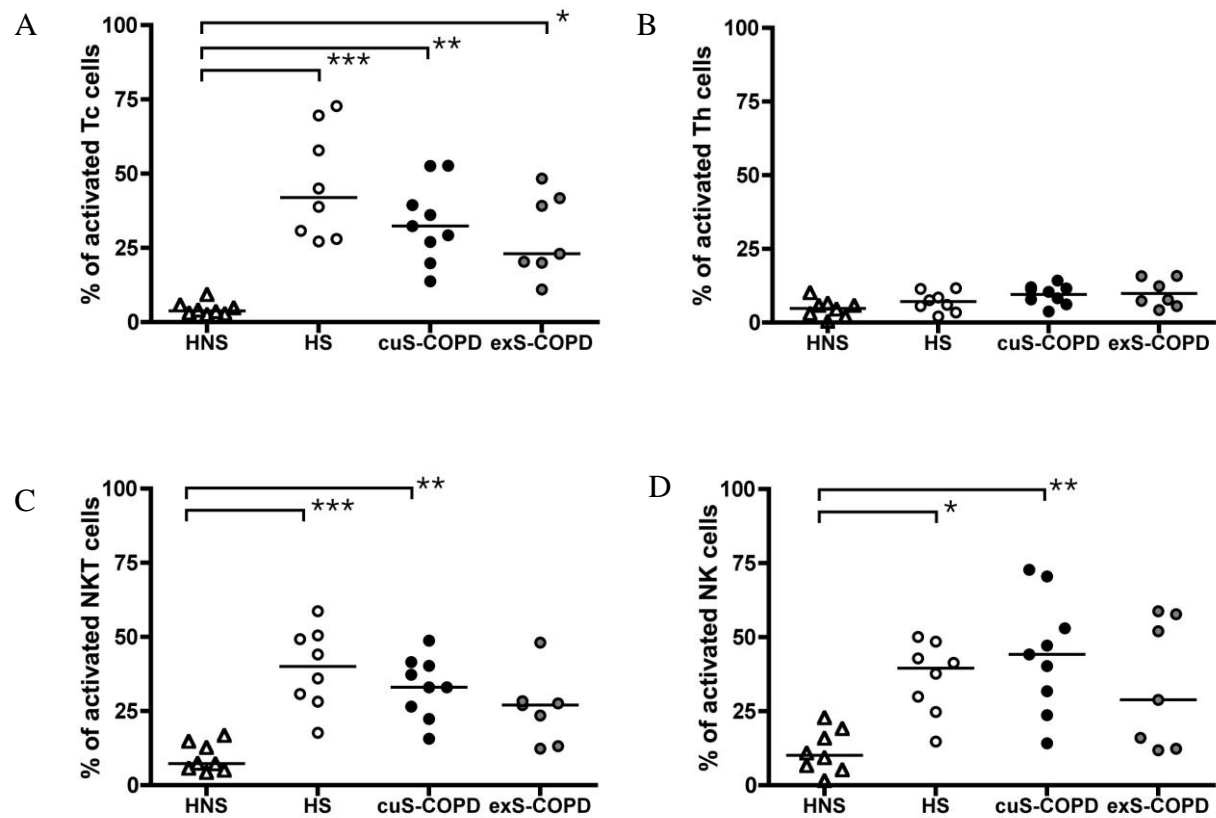


Figure 3.1.4 Activation of killer cell types from peripheral blood *ex vivo* of four groups. Activation of CD8⁺ T cells, CD4⁺ T cells, NKT-like cells and NK cells were analysed. There was a significant increase in level of activation of CD8⁺ T cell (panel A), NKT-like cells (panel C), and NK cells (panel D) from current healthy smokers (HS, open circles) and current smokers with COPD (cuS-COPD, black solid circles). Activation of NK cells from ex-smokers with COPD (exS-COPD, grey solid circles) was also significantly increased compared to healthy non-smokers (HNS, open triangle). There was no statistical difference in activation of CD4⁺ T cells (panel B) between groups. *: $p < 0.05$, **: $p < 0.01$, ***: $p < 0.001$. HNS (n=8), HS (n=8), cuS-COPD (n=9), and exS-COPD (n=7).

activated *ex vivo* than those from healthy non-smokers (10.1%, 1.5-22.8) (Figure 3.1.4.D). There were no significant differences in NK cell activation between ex-smokers with COPD (28.9%, 11.91-58.8; $p>0.05$) and other groups of interest.

Interestingly, few $CD4^+$ T lymphocytes were activated in peripheral blood from four groups (Figure 3.1.4.B) and there were no statistical differences between groups (all p value >0.05), namely, healthy non-smokers (5.17%, 0.4-10.16), current healthy smokers (6.87%, 2.16-11.7), current smokers with COPD (10.46%, 3.81-14.31) and ex-smokers with COPD (7.81%, 4.24-15.89).

3.1.3.3 Expression of CD25 on $CD4^+$ T cells

Although activation of $CD4^+$ T lymphocytes was lower than other cell types of interest (Figure 3.1.4.B), activation markers including CD69 and CD25 were still detectable on their cell surface (Figure 3.1.3.B). Regulatory T (Treg) cells, as a subset of $CD4^+$ T cells, also express CD25. Therefore, we hypothesised that those $CD4^+CD25^+$ T cells could be Treg cells. To examine this further we generated dot plots of Forward Scatter (FS) vs Side Scatter (SS) that showed, unlike other activated cell types expressing CD25 that were larger and

more granular (Figure 3.1.5.A, C and D), those CD4⁺ CD25⁺ T cells were primarily small and less granular and thus quite probably inactivated lymphocytes (Figure 3.1.5.B). These observations were seen in all groups of interest.

3.1.3.4 Correlation of cell activation and number of cigarettes being smoked per day

In view of our findings that current smoking increased numbers of activated Tc, NKT and NK cells, we looked for correlations between the number of cigarettes currently being smoked per day and level of *ex vivo* cell activation for these three cell types from both current healthy smokers and current smokers with COPD (Figure 3.1.6). Cells expressing CD69 or CD25 alone, and expressing both activation markers were considered as activated cells in each cell type. In current smokers with COPD, NK cells showed a significant positive correlation between the proportion of *ex vivo* activation and number of cigarettes smoked ($r=0.7615$, $p=0.0214$) (Figures 3.1.6.C). No such correlation was observed for CD8⁺ T cells ($r=0.2427$, $p=0.5206$) (Figure 3.1.6.A), NKT-like cells ($r=0.0753$, $p=0.8432$) (Figure 3.1.6.B). Also, there was no significant correlation between the proportion of *ex vivo* activation and number of cigarettes smoked in these cell types from current healthy smokers (Figure 3.1.6).

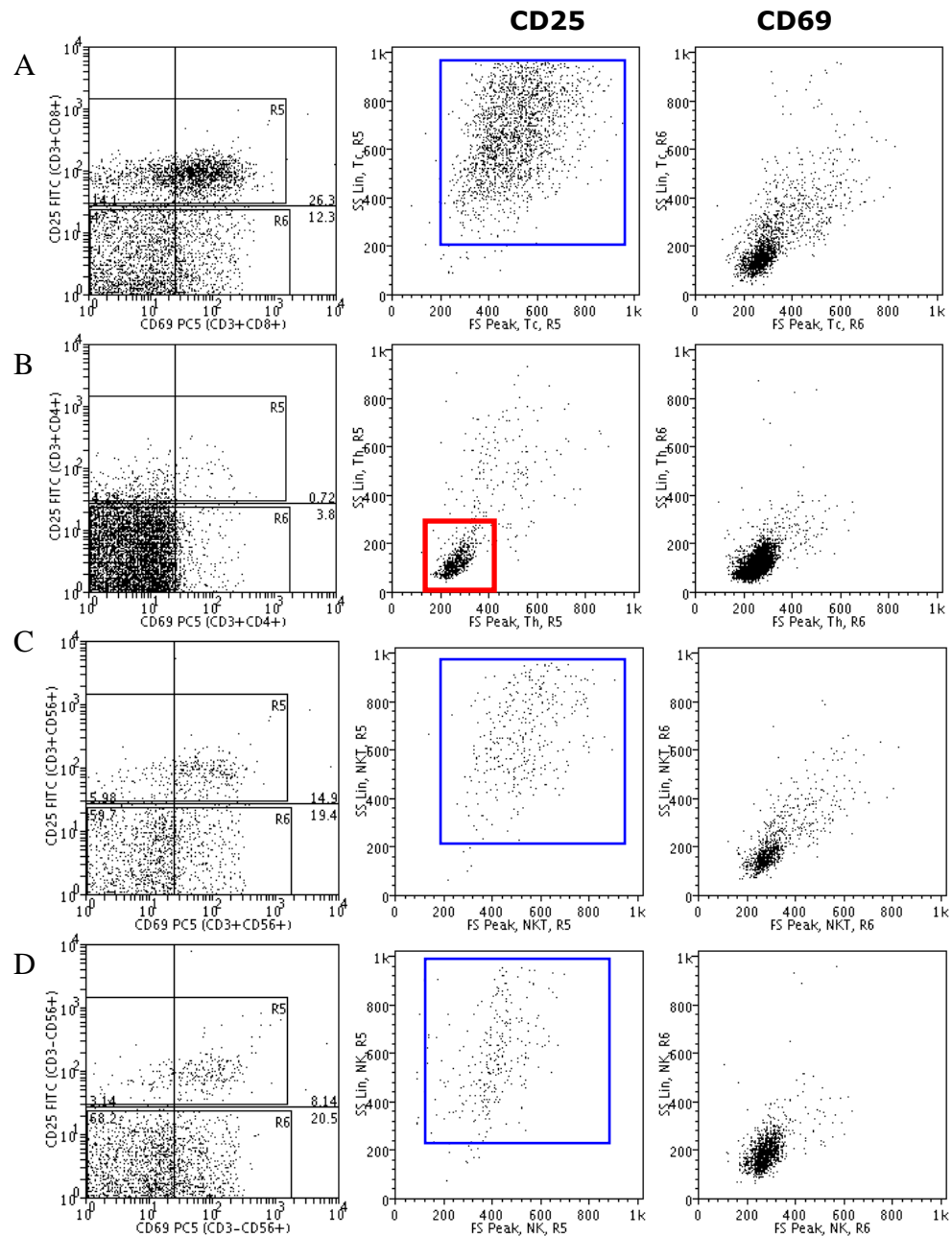


Figure 3.1.5 Representative dot plots for gating cell types that express CD25 from current smokers with COPD. CD69 and CD25 are cell surface markers of early and later stages of lymphocyte activation, respectively. Unlike other cell types of interest, CD4⁺ T cells expressing CD25 was mainly lymphocytes (red square) but not activated cells (blue square). Regulatory T (Treg) cells are a subset of CD4⁺ T cells expressing CD25, which indicated that those CD4⁺CD25⁺ T cells might be Treg cells. (A) CD8⁺ T cell; (B) CD4⁺ T cells; (C) NKT-like cells; and (D) NK cells.

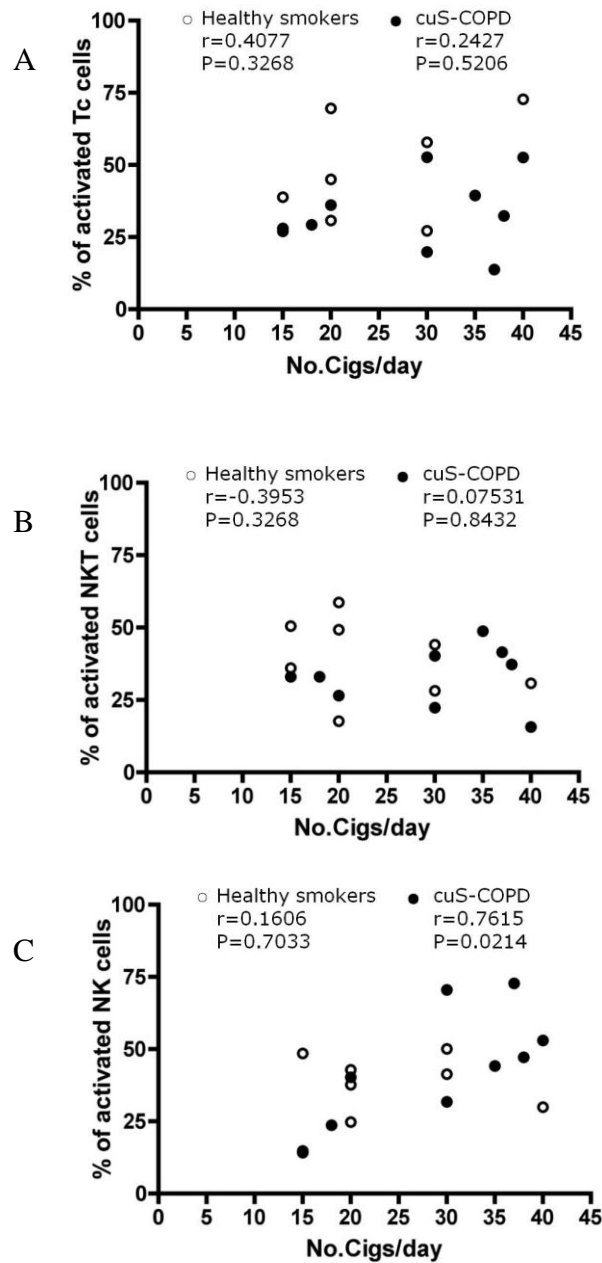


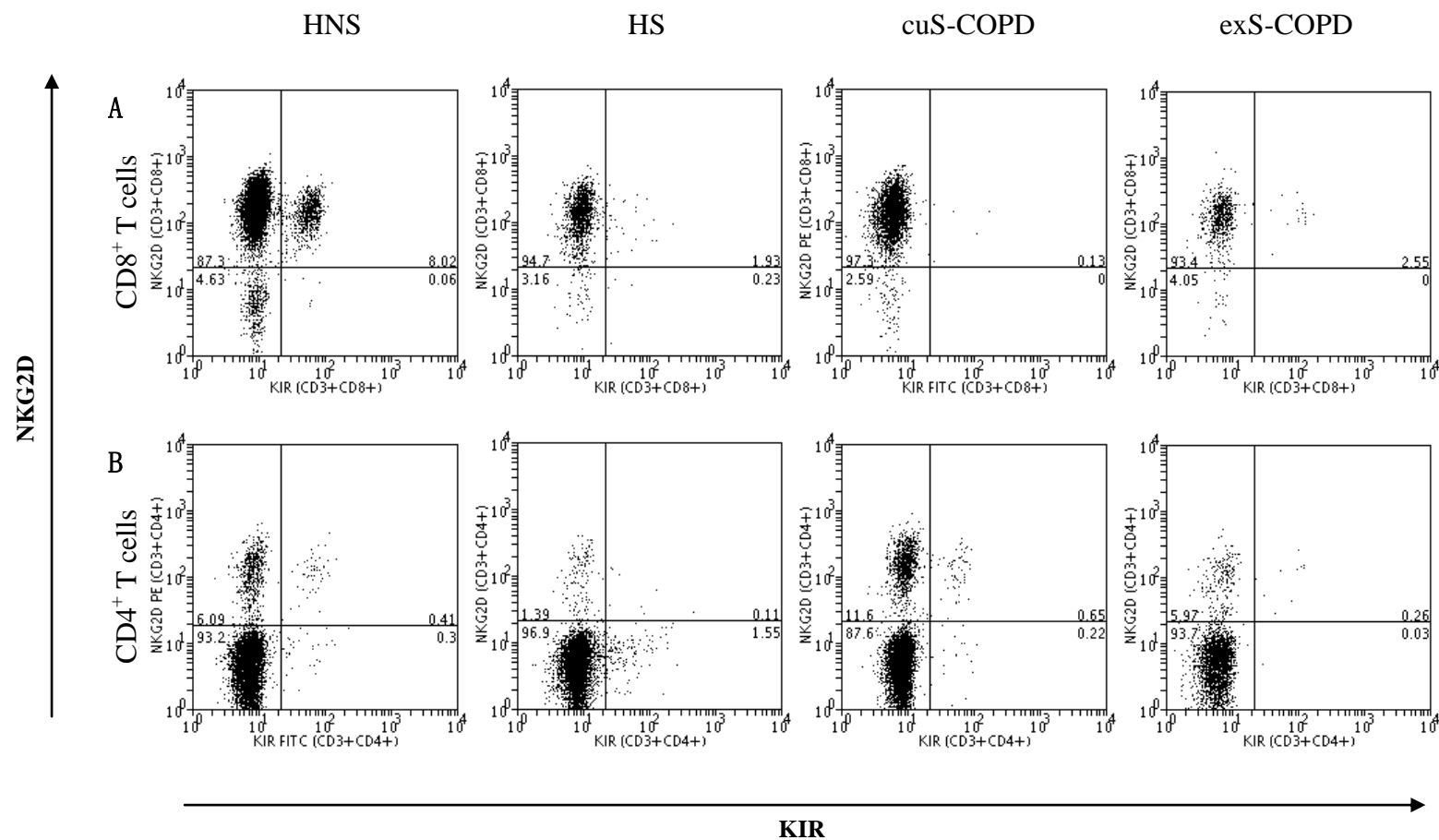
Figure 3.1.6 Correlation of the proportion of cells activated and number of cigarettes currently smoked per day in current healthy smokers (HS) and current smokers with COPD (cuS-COPD). Analysis was performed on peripheral blood CD8⁺ T lymphocytes (Panel A), NKT-like cells (Panel B) and NK cells (Panel C). A significant correlation was observed in NK cells from current smokers with COPD. HS, n=8; cuS-COPD, n=9.

3.1.3.5 KIR (CD158e1) and NKG2D expression in peripheral blood

The balance between activation receptors and inhibitory receptors expressed by cells is one of crucial factors orchestrating cell activation. Expression of both the inhibitory KIR (CD158e1) and the activating receptor NKG2D by killer cells, including CD8⁺ T cells (Figure 3.1.7.A), CD4⁺ T cells (Figure 3.1.7.B), NKT cells (Figure 3.1.7.C) and NK cells (Figure 3.1.7.D), were measured by flow cytometry. Representative dot plots of KIR and NKG2D expression from healthy non-smokers (HNS); current healthy smokers (HS); current smokers with COPD (cuS-COPD); and ex-smokers with COPD (exS-COPD) are shown in Figure 3.1.7.

CD8⁺ T-lymphocytes showed a significantly lower proportion of cells from healthy smokers (0.83%, 0.12-2.16; $p < 0.001$) and current smokers with COPD (0.66%, 0.13-3.97; $p < 0.001$) expressed CD158e1 on the cell surface compared to healthy non-smokers (9.91%, 4.78-16.6) (Figure 3.1.8.A).

The proportion of NKT-like cells expressing surface CD158e1 was also significantly lower in current healthy smokers (2.14%, 0.24-7.92; $p < 0.001$) and current smokers with COPD (2.24%, 1.58-6.77; $p < 0.001$) compared to healthy



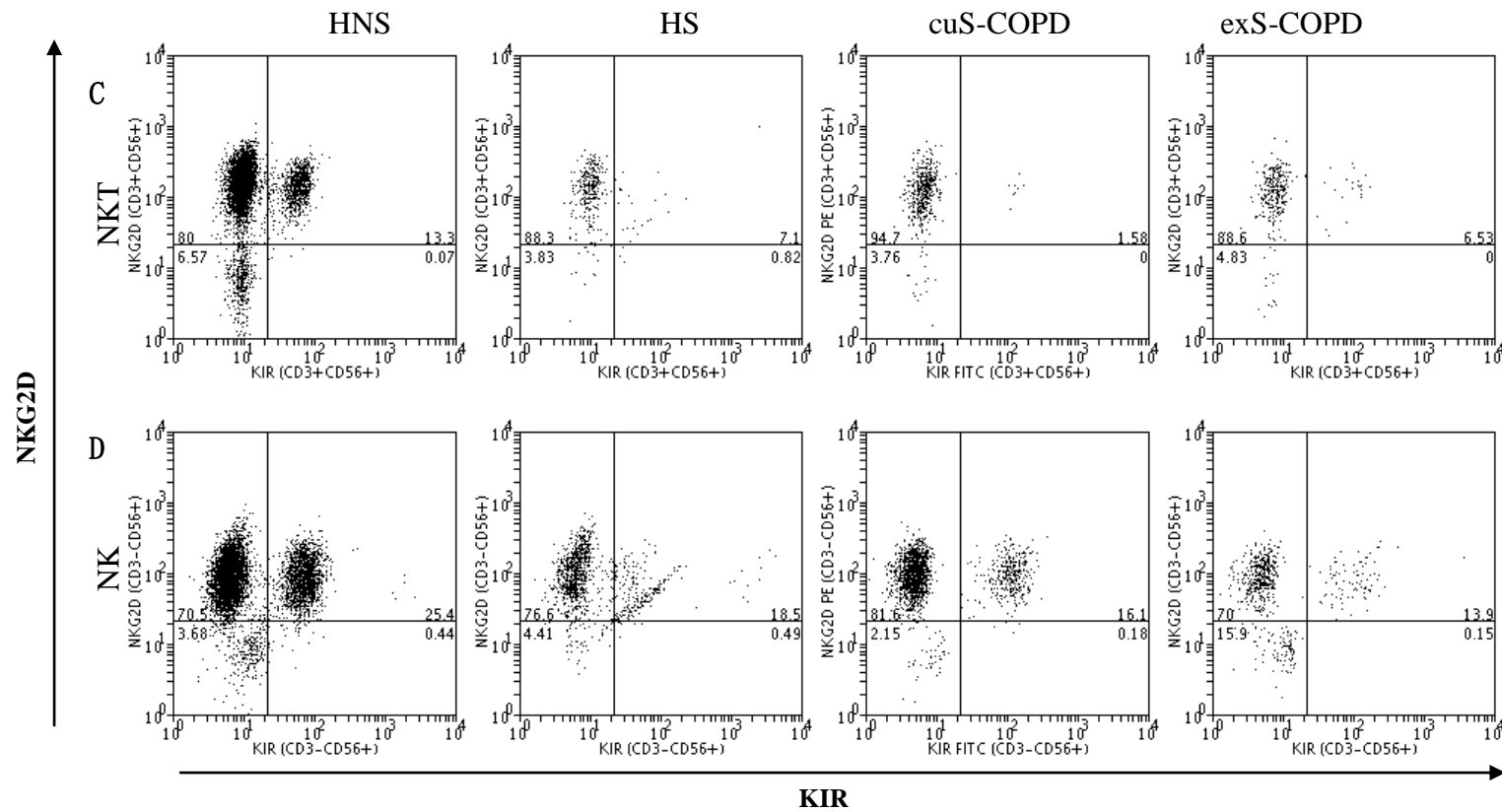


Figure 3.1.7 Representative dot plots for KIR (CD158e1) and NKG2D expression of cell types of interest from four groups including healthy non-smokers (HNS), current healthy smokers (HS), current smokers with COPD (cuS-COPD) and ex-smokers with COPD (exS-COPD). NKG2D was highly expressed by CD8⁺ T cells, NKT-like cells and NK cells but not CD4⁺ T cells. (A) CD8⁺ T cells; (B) CD4⁺ T cells; (C) NKT-like cells; and (D) NK cells.

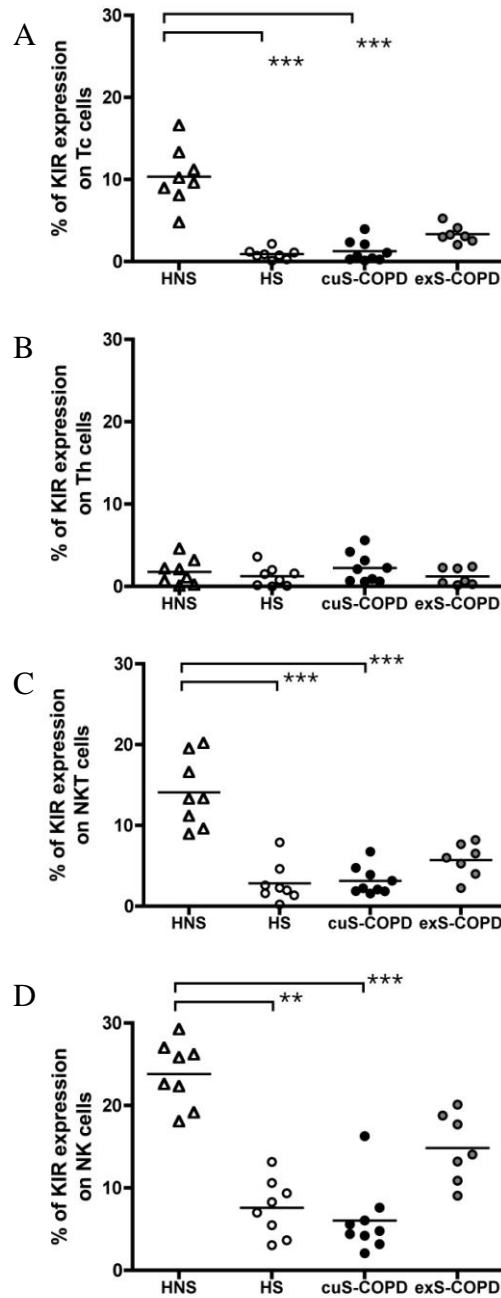


Figure 3.1.8 KIR (CD158e1) expression of cell types of interest from peripheral blood *ex vivo* of four groups. Levels of KIR expression by CD8⁺ T cell (panel A), NKT-like cells (panel C), and NK cells (panel D) were all significantly reduced in current healthy smokers (HS, open circles) and current smokers with COPD (cuS-COPD, solid circles) compared to healthy non-smokers (HNS, open triangles). There was no significant difference in the KIR expression between ex-smokers with COPD (exS-COPD, grey solid circles) and other groups. There was also no statistical difference in the levels of KIR expression on CD4⁺ T cells (panel B) between groups. *: $p < 0.05$, **: $p < 0.01$, ***: $p < 0.001$. HNS (n=8), HS (n=8), cuS-COPD (n=9), and exS-COPD (n=7).

non- smokers (13.34%, 8.97-20.2) (Figure 3.1.8.C).

Analysis of NK cells showed the proportion of these cells expressing surface CD158e1 was also significantly lower in current healthy smokers (7.7%, 3.06-13.17; $p < 0.01$) and current smokers with COPD (4.79%, 2.08-16.28; $p < 0.001$) compared to healthy non-smokers (24.22%, 18.1-29.26) (Figure 3.1.8.D).

No differences in proportions of CD4⁺ T-lymphocytes (in all cases median levels <2.1%) expressing CD158e1 were shown between the four groups (Figure 3.1.8.B).

Interestingly, proportions of CD4⁺ T lymphocytes expressing NKG2D were significantly lower (in all cases median levels <9.0%) compared to proportions of CD8⁺ T-lymphocytes, NKT-like cells and NK cells (in all cases over 95%) in each group. Meanwhile, there were no statistical differences on the proportion of each cell type expressing NKG2D between four groups (Figure 3.1.9).

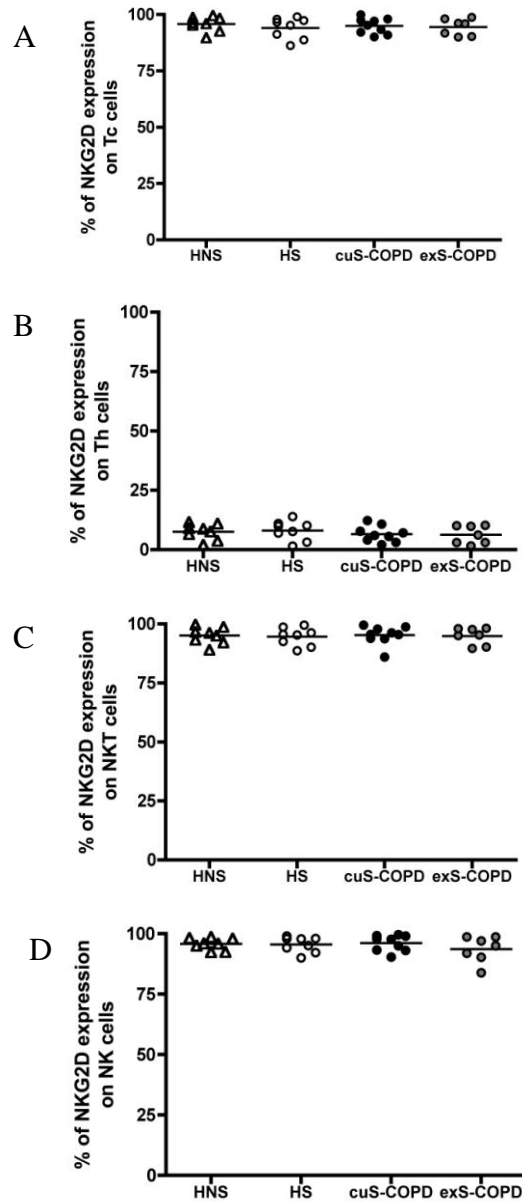


Figure 3.1.9 Proportions of NKG2D expression on cell types of interest from peripheral blood *ex vivo* of four groups. NKG2D expression by CD8⁺ T cells (Tc, panel A), CD4⁺ T cells (Th, panel B), NKT-like (NKT, panel C) cells and NK cells (NK, panel D) were analysed. There were no significant differences on the proportion of NKG2D expression by each cell type from four groups, including healthy non-smokers (n=8), current healthy smokers (n=8), current smokers with COPD (cuS-COPD, n=9), and ex-smokers with COPD (exS-COPD, n=7). All p value >0.05.

3.1.4 Discussion

Here we report, for the first time, that current smoking *per se* is associated with a significantly higher proportion of peripheral blood cytotoxic cells that are activated *ex vivo* and this is positively correlated with the number of cigarettes currently smoked per day.

We and others have previously demonstrated the potential involvement of the three main classes of human killer cells, namely, CD8⁺ T-lymphocytes, NKT-like cells and NK cells in COPD (Paats et al., 2012, Roos-Engstrand et al., 2009, Roos-Engstrand et al., 2010, Urbanowicz et al., 2009). These killer cells play key roles in inflammatory responses and activation of these cells can cause the production of inflammatory cytokines and chemokines, such as TNF- α and IL-8, that can induce pathological features of COPD and lung tissue damage (Barnes, 2004c). Therefore, we hypothesized that killer cells are more active in COPD patients. Here we examined the *ex vivo* activation phenotype of these cells in the peripheral blood. In peripheral blood, current smokers with or without COPD, but not ex-smokers with COPD, have the highest proportion of activated CD8⁺ T-lymphocytes, NKT-like cells and NK cells compared with healthy non-smokers and ex-smokers with COPD. This suggests that smoking *per se* affects systemic

killer cell activation. We thus examined the correlation between current smoking profiles and cell activation levels. Here we show that NK cell activation is positively correlated with the number of cigarettes currently smoked in COPD patients. No correlations between cell activation and lung function (FEV₁ percentage of predicted) were found (data not shown). These data suggest that in peripheral blood, cell activation levels are more related to current smoking habit rather than disease. A previous study examining the effect of smoking cessation has shown in BAL fluid that CD8⁺ T-lymphocytes from both smokers with normal lung function and COPD participants expressed high levels of CD69 and CD25 activation markers (Roos-Engstrand et al., 2009). Sputum T-lymphocytes have also been shown to express higher levels of CD103 and CD69 than blood lymphocytes suggesting higher numbers of activated intraepithelial phenotype T-lymphocytes in the lung (Leckie et al., 2003). Furthermore, activated CD8⁺ T cells secrete more CXCR3 that lead to the accumulation of CD8⁺ T cells to the infected tissue and organs, which may explain why CD8⁺ T cells are predominant increased in COPD but not CD4⁺ T cells. Interestingly, here we also show few CD4⁺ T cells are activated in peripheral blood from the four groups. We further show that those CD4⁺CD25⁺ T cells may be regulatory T cells (not activated cells) by dot plot of

Forward Scatter (FS) vs Side Scatter (SS). To confirm this observation further studies are required. Although two studies have found highly activated Th1 cells in the lung of ex-smokers with COPD and in the small airway wall of smokers with severe COPD by detecting their chemokine and cytokine secretion, those samples are actually from COPD patients with emphysema or co-suffering from cancer (Turato et al., 2002).

One of the crucial factors determining cell activation is the signaling balance between activating receptors and inhibitory receptors. To investigate a possible mechanism for differences in activation states observed in peripheral blood, we compared expression of the inhibitory receptor KIR CD158e1 and the activating receptor NKG2D. Interestingly, in this study, the expression levels of the inhibitory KIR CD158e1 were lower in peripheral blood of current smokers with or without COPD compared to healthy non-smokers, which was the opposite to the activation levels of these killer cells including CD8⁺ T cells, NKT-like cells and NK cells. Similar study supports this finding. CD94 is another inhibitory receptor expressed by NKT-like cells and NK cells (Lazetic et al., 1996). Decreased expression of CD94 on both NKT-like cells and NK cells were detected in PBMCs from COPD patients compared to healthy volunteers (Hodge et al., 2012). However, they did not detect any

significant differences on the cell activation levels of NKT-like and NK cells (only CD69 expression) between groups. Therefore, our findings provided the first demonstration of association of cell activation and KIR expression. We also show that activating receptor NKG2D are highly expressed by CD8⁺ T cells, NKT-like cells and NK cells but not CD4⁺ T cells, which is in accordance with previous studies that demonstrate NKG2D receptors are constitutively expressed and almost exclusively on cytotoxic cells including CD8⁺ T cells, NKT cells, NK cells and $\gamma\delta$ T cells (Maasho et al., 2005). There are no significant differences of NKG2D expression on CD8⁺ T cells, CD4⁺ T cells, NKT-like cells and NK cells respectively, in the four groups. This further supports the proposal of increased peripheral blood killer cell activity in smokers due to the increased ratio of activating to inhibitory receptors.

The present study does have some limitations that deserve comment. Firstly, we used CD69 and CD25 as activation markers. Although other markers exist, these are generally accepted as representing cell activation (Marquez et al., 2010, McDonald et al., 2010). Secondly, CD25 is commonly used as a marker for T regulatory cells (Lane et al., 2010). To further confirm whether these CD4⁺CD25⁺ T cells are Treg cell, FOXP3 can be used as a good marker for Treg cell identification.

Also, fluorophore FITC is not particularly bright, they photobleach rapidly and are self-quenching at high concentration. Therefore, other fluorophore, such as the AlexaFluors, may be applied in future study to gain high quality analysis. Thirdly, there are many kinds of inhibitory and activating receptors expressed on these cells but due to limited cell numbers we analysed two representative receptors that give an indication of the killing function of these cells. Fourthly, it should be noted that the majority of participants with COPD were on inhaled corticosteroids. However, the two participants who were not on inhaled corticosteroids in this group showed the same results in terms of activation and KIR expression as those on inhaled corticosteroids.

Of the note, our colleague also detected, *ex vivo*, the activation levels of killer cells in sputum samples from COPD patients (data shown in the publication paper). Unlike systemic activation, lung activation is due specifically to COPD, irrespective of current smoking status.

In summary, we have shown for the first time the activation levels of killer cells from peripheral blood of COPD patients. Also, the systemic activation is in response to smoking *per se*. We further demonstrate expression levels of inhibitory

receptor KIR (CD158e) in COPD groups. These findings indicate that activation of killer cells in COPD participants may play a role in the pathogenesis of COPD.

3.2 Effects of cigarette smoke extract on killer cell activation in COPD

3.2.1 Introduction

Cigarette smoke extract (CSE)

Cigarette smoke has been identified as the main risk factor in COPD development. Previous studies have demonstrated that cigarette smoke affects structural and functional changes in the respiratory ciliary epithelium (Cantin et al., 2006, Mio et al., 1997, Wyatt et al., 1999), lung surfactant protein and immune cells (Mohan, 2002a), such as lymphocytes, alveolar macrophages, dendritic cells, neutrophils and NK cells. Therefore, accumulating data suggested that chronic inhalation of cigarette smoke alters a wide range of immunologic function, including both innate and adaptive immune responses.

The smoke from a cigarette contains thousands chemicals which have various toxic, mutagenic and carcinogenic activity (Hoffmann and Wynder, 1986, Stedman, 1968). The concentrations of chemical ingredients also vary widely in the different cigarette brands. Studies show that various components of cigarette smoke can affect the immune

response in different cell types (Lee et al., 2007b, McCue et al., 2000, Ouyang et al., 2000). Two major components of smoke are nicotine and tar (Hoffmann and Wynder, 1986, Stedman, 1968). Previous studies have shown that nicotine is tightly associated with increased risk of cardiovascular disease and cancer, particularly of lungs. Tar is an irritant. It enters the respiratory system as an aerosol of minute droplets that are deposited in the respiratory system causing excessive mucus production. Tar also contains carcinogens that combine with the affected mucus membranes. It causes lung cancer, chronic bronchitis and emphysema. Holt *et al* has investigated that cigarette smoke containing high levels of tar and nicotine is more immunosuppressive than the smoke from low-tar low-nicotine cigarettes (Holt et al., 1976). However, effects of cigarette smoke on the immune system are biphasic, which is mainly determined by the components and duration of exposure. For example, unlike nicotine and tar, tobacco glycoprotein and metals present in cigarette smoke are generally immunostimulatory (Brooks et al., 1990, Francus et al., 1992)..

Immunological features of cigarette smoke in COPD have been explored in human studies. *In vitro* activation of T cells and expression of granzyme B were significantly suppressed by

cigarette smoke, which resulted in altered immune response of the lymphocytes (Glader et al., 2006a). Skold *et al* have demonstrated that exposure to cigarette smoke altered the phenotype of human alveolar macrophages (Skold et al., 1996). Recently, effects of cigarette smoke on DCs in COPD have been more explored. *In vitro*, the effects of cigarette smoke on DCs were dose-dependent. Production of cytokines such as IL-12 and IL-10 during DCs maturation process can influence DC induction of a Th1 or Th2 immune response. In addition to expressing high levels of antigen-presenting molecules and costimulatory molecules, mature DCs release large amounts of IL-12 that can stimulate a Th1 immune response. Release of IL-10, however, blocks the DC maturation process by interfering with up-regulation of costimulatory molecules and production of IL-12, subsequently limiting the ability of DCs to initiate a Th1 response (Corinti et al., 2001, De Smedt et al., 1997). Early studies have shown that Low doses of nicotine enhanced human monocyte-derived DC maturation and their capacity to secrete IL-12 (Aicher et al., 2003). On the contrary, higher doses of nicotine treated DCs had low endocytic and phagocytic function, produced less IL-12 and more Th2 cytokine IL-10 that promoted DCs priming Th2 cell differentiation more efficiently (Nouri-Shirazi and Guinet, 2003,

Nouri-Shirazi and Guinet, 2006). Similar results were also shown in Vassallo *et al* study. LPS-induced DC maturation was suppressed by cigarette smoke, while the secretion of IL-10 and prostaglandin E2 from these DCs were enhanced (Vassallo *et al.*, 2005). Moreover, cigarette smoke impaired the production of TNF- α , IL-6 and IFN- α , whereas augmented the IL-8 secretion by plasmacytoid dendritic cells in response to Toll-like receptor (TLR)-9 (Mortaz *et al.*, 2009). Studies on the effects of cigarette smoke on NK cell properties showed controversial findings. Some groups found that cigarette smoke impaired NK cell cytotoxic activities (Takeuchi *et al.*, 1988), whereas others showed that their killer functions were increased or not affected (Meliska *et al.*, 1995, Newman *et al.*, 1991). However, these findings were generated from different disease models. Co-activation of NK cells by cigarette smoke and NK cell inducers have been established in Mian *et al* studies. Cigarette smoke greatly impaired the production of IFN- γ and TNF- α by NK cells upon polyinosinic:polycytidylic acid (poly I:C) activation. The cytotoxic activities and perforin expression of poly I:C activated NK cells were also attenuated by cigarette smoke (Mian *et al.*, 2008). Treatment of neutrophils with cigarette smoke extract can suppress neutrophil caspase-3-like activity and the phagocytic uptake

and intracellular digestion of staphylococci by neutrophils (Stringer et al., 2007).

IL-2 and IL-15

More than 30 years ago, IL-2 was shown to be a T cell growth factor purified from mitogen-stimulated lymphocyte culture (Morgan et al., 1976). Twenty years later, IL-15 was also identified through its ability to mimic IL-2-induced T cell proliferation (Bamford et al., 1996, Grabstein et al., 1994). Both cytokines belong to the 4- α -helix bundle cytokine family and they share several signaling pathways through their heterotrimeric receptors (Lin and Leonard, 1997, Lin et al., 1995, Miyazaki et al., 1994, Fujii et al., 1995, Kawahara et al., 1995, Waldmann and Tagaya, 1999). Receptors for IL-2 (IL-2R) and IL-15 (IL-15R) are comprised of three subunits, namely, IL-2R α (IL-15R α for IL-15), IL-2/15R β (also known as CD122) and common cytokine-receptor γ -chain (γ c) (Anderson et al., 1995, Asao et al., 2001, Cosman et al., 1995). To achieve high affinity form of receptor, three subunits must be presented together. In the absence of IL-2R α or IL-15R α , receptors for IL-2 or IL-15 will only show intermediate-affinity. Signal inductions are relied on β and γ chains of receptors (Minami et al., 1993, Smith, 1988a, Smith, 1988b, Waldmann et al., 1998). IL-2R α and IL-15R α are two unique subunits for

IL-2 and IL-15 respectively with also distinct physiologic expression profiles. IL-2R α binds to IL-2 with low-affinity without signal transduction, they are only detectable on activated lymphocytes and Treg cells (Fontenot et al., 2005, Rogers et al., 1997). Whereas IL-15R α binds to IL-15 with high-affinity and recent studies have shown that IL-15R α plays functions on intracellular signals through binding to tumour-necrosis factor-receptor-associated factor 2 (TRAF2) through its cytoplasmic tail and preventing interaction between IL-15R α and the pro-apoptotic cascade members (Bulfone-Paus et al., 1999, Pereno et al., 1999, Pereno et al., 2000). IL-15R α has a wide cellular distribution. It can be observed in multiple tissues and cell types, such as in liver, heart, spleen, lung and activated vascular endothelial cells (Giri et al., 1995), T cells, B cells, activated monocytes, dendritic cells, macrophages and bone marrow stromal cell lines (Anderson et al., 1995). Distribution of three subunits of IL-2R and IL-15R, namely, α -chain, β -chain (CD122) and γ -chain, is shown in Table 3.2.1.

Table 3.2.1 Expression of three units of IL-2R and IL-15R on immune cells

Cell types	IL-2Rα	IL-15Rα	IL-2/15Rβ	γc
Pre-B cells	+	+	-	+
Mature B cells	-	+	-	+
Mature T cells				
Naïve	-	+	+	+
Effector	+	+	+	+
Memory	-	+	+	+
Treg cells	+	-	+	+
NK cells	-	+	+	+
Dendritic cells	+/- *	+	+/-*	+
Monocytes	-	+	-	+

*Certain activated dendritic cells have been shown to express IL-2R α and IL-2/15R β on their surfaces (Ardavin and Shortman, 1992, Mnasria et al., 2008).

Due to common features in receptor components and signaling pathways, IL-2 and IL-15 have several similar functions. They can induce the proliferation of activated T cells, generate cytotoxic T cells and NK cells and facilitate B cell proliferation and immunoglobulin synthesis (Fehniger and Caligiuri, 2001, Waldmann et al., 2001, Waldmann and Tagaya, 1999). However, in many adaptive immune responses, significant differences between IL-2 and IL-15 have been identified in their molecular and cellular biology and functions (Thomas, 2008, Waldmann and Tagaya, 1999). The distinct binding affinity and distribution of two α -chains may be one reason contributing to distinct features of two receptors. As mentioned previously, IL-2 α is mainly expressed by activated

T cells and bind IL-2 with low affinity, while IL-15 α binding to IL-2 with high affinity is widely spread in many tissues and cell types. Also, in contrast to IL-2 that binds pre-formed high affinity heterotrimeric receptors at the cell surface (Taniguchi and Minami, 1993, Waldmann, 1991), IL-15 is only secreted at low levels and binds membrane IL-15R (Dubois et al., 2002). Functional differences may result in distinctive roles of IL-2 and IL-15 in human diseases. For example, IL-2 can down-regulate γ chain expression on cycling T cells and dampen T cell clonal expansion through activation-induced cell death (AICD) (Lenardo, 1991), whereas IL-15 is an anti-apoptosis factor and inhibits IL-2 induced AICD by facilitating the maintenance of CD8⁺ T cell survival, including self-reactive memory T cells (Marks-Konczalik et al., 2000). Furthermore, IL-2 participates in the maintenance of peripheral Treg cells (D'Cruz and Klein, 2005, Fontenot et al., 2005, Maloy and Powrie, 2005) but IL-15 has no marked effects on them (Fehniger and Caligiuri, 2001a). IL-2 shows ability of inhibiting CD8⁺ memory T cell proliferation but IL-15 can selectively stimulated the proliferation of memory CD8⁺ T cells and prolong their survival (Zhang et al., 1998). Collectively, although both IL-2 and IL-15 play potent roles in the pathogenesis of autoimmune diseases, cancers and inflammatory diseases, they can be either beneficial of

Table 3.2.2 Comparison of human IL-2 and IL-15

Properties	IL-2	IL-15
Structure	15.5kDa; four-helix bundle	14-15kDa; four-helix bundle
Receptors	IL-2R α , low affinity; IL-2/15R β ; γ C	IL-15R α , high affinity; IL-2/15R β ; γ C
Producing cells	Activated T cells	Monocytes, dendritic cells and epithelial cells*
Function	Proliferation of T cells, NKT cells, NK cells and B cells; Activating NK cells and NKT cells; Elimination of self-reactive T cells; Inducing AICD; Maintenance of Treg Cells	Proliferation and activation of NKT cells and NK cells; Maintenance of memory CD8 ⁺ T cells and NK cell survival; Inhibiting IL-2-induced AICD

* Epithelial cells are big producers of IL-15. Respiratory viruses such as respiratory syncytial virus (RSV) increased surface MIC molecules in respiratory epithelial cells (Zdrengea et al., 2012). These molecules are ligands for NKG2D receptors on NK and T cells, playing an important role in the nonspecific activation of cytotoxic activity of NKG2D-positive NK and CD8⁺ T cells.

detrimental. Their application in immune therapy may also differ at some points due to their unique biological features. Table 3.2.2 simply outlines the comparison of human IL-2 and IL-15.

To date, IL-2 and IL-15 biological functions and their therapeutic applications have been well studied in cancers, viral infections and certain autoimmune diseases.

Nevertheless, their properties in smokers or COPD patients are still less explored. Higher IL-2 levels have been found in lung plasma samples from stable COPD patients compared to participants with rapidly progressive COPD (D'Armiento et al., 2009). Also, corticosteroid (dexamethasone) have shown less inhibition of IL-2 production in COPD patients compared to healthy volunteers although these numerical differences between groups did not reach statistical significance (Kaur et al., 2012). *In vitro*, it has shown that IL-15 stimulation by itself had no effect on IFN- γ protein expression in COPD patients at GOLD stages 2, 3, and 4. However, IL-15 primed CD8⁺ lung T cells to have greater cytotoxic potential by expressing higher levels of intracellular perforin when they encounter a target cells in COPD (Freeman et al., 2010). Furthermore, CSE treated NK cells were less activated after poly I:C stimulation, and this suppression was IL-15 mediated (Mian et al., 2009).

Accumulating evidence strongly suggest CSE has markedly association with COPD developments. Although numbers of studies have shown that CSE impairs cytokine and chemokine productions in many cell types, for instance, macrophages, neutrophils, DCs or other inflammatory cells in healthy volunteers (Gaschler et al., 2008, Takahashi et al., 2007,

Takahashi et al., 2006), however, the impact of CSE on killer cells from COPD patients remains to be explored. Previous section has shown that killer cells were all significantly activated *in vivo* in current smokers with or without COPD, which indicates that cigarette smoking, rather than disease itself, may play a potent role in killer cell activation in peripheral blood. α -CD3 is a great stimulator for CD3⁺ T cell activation. Whereas, IL-2 and IL-15 can induce NKT cells and NK cell stimulation. In this chapter, we have investigated, *in vitro*, the effects of CSE on unstimulated or α -CD3, IL-2 or IL-15 activated killer cells including CD8⁺ T cells, NKT-like cells and NK cells, from peripheral blood in four designed groups, namely, healthy non-smokers (HNS), current healthy smokers (HS), current smokers with COPD (cuS-COPD) and ex-smokers with COPD (exS-COPD).

3.2.2 Material and methods

3.2.2.1 Study population and procedures

The criteria of participant enrollments are the same as previously mentioned. Twenty participants were included in CSE activation study. Table 3.2.3 and Table 3.2.4 detail the demographic and spirometric data of the participants in

Table 3.2.3 Demographic and spirometric value of the healthy non-smoking participants in optimal concentration experiments

Healthy non-smokers	Age (years)	Gender (M/F)	FEV ₁ (% pred)	FEV ₁ /FVC (%)	ΔFEV ₁ post bronch	BMI (kg/m ²)
n=4	49 (43-57)	3/1	110 (89-118)	99 (82-111)	2.3 (2.0-2.8)	22.8 (19.6-27.7)

Table 3.2.4 Demographic and spirometric value of the CSE suppression study groups

	Healthy non-smokers	Healthy smokers	cuS-COPD	exS-COPD
Participants	4	4	4	4
Age (years)	55 (53-60)	54 (50-58)	64 (55-73)	56 (49-65)
Gender (M/F)	1/3	3/1	3/1	2/2
Packs/yr	0	36 (32-40)	47 (42-66)	34 (23-47)
Smoking status (Current/Ex)	0	4/0	4/0	0/4
Chronic bronchitis (Yes/No)	0	2/2	3/1	1/3
FEV ₁ (% pred)	103 (81-116)	98 (89-106)	43 (37-52)	59 (54-63)
FEV ₁ /FVC (%)	94 (76-108)	83 (70-99)	36 (32-46)	52 (37-57)
ΔFEV ₁ post bronch	2.3 (2.0-2.6)	2.0 (1.9-2.4)	1.6 (1.5-2.2)	1.2 (0.9-1.3)
BMI (kg/m ²)	23.6 (20.7-26.0)	26.0 (21.8-28.5)	25.9 (18.9-29.8)	21.1 (20.0-22.7)
Inhaled GCS (on/off)	N/A	N/A	1/3	1/3
MRC dyspnoea scale	N/A	N/A	3 (2-4)	3 (2-4)
Distance walked in 6 min (m)	N/A	N/A	268 (102-364)	336 (152-428)
BODE Index	N/A	N/A	5 (3-6)	4 (2-5)

preliminary concentration experiments and CSE suppression study respectively.

Table 3.2.5 Antibodies used for flow cytometry

Antigen	Fluorochrome	Isotype	Clone	Source
CD3	PC7	Mouse IgG1	UCHT1	Beckman Coulter, Luton, UK
CD8	ECD	Mouse IgG1	SFCl21Th y2D3	Beckman Coulter, Luton, UK
CD25	FITC	Mouse IgG2a,	B1.49.9	Beckman Coulter, Luton, UK
CD56	PE	Mouse IgG1	N901	Beckman Coulter, Luton, UK
CD69	PC5	Mouse IgG2b	TP1.55.3	Beckman Coulter, Luton, UK

3.2.2.2 Reagents

Anti-CD3 was purchased from R&D system (Abingdon, UK). Recombined human IL-2 (rhIL-2) and recombined human IL-15 (rhIL-15) were obtained from PeproTech EC (London, UK). Other monoclonal antibodies for flowcytometry were shown in table 3.2.5.

3.2.2.3 Peripheral Blood Mononuclear Cell isolation

PPBMCs were isolated from whole blood on a discontinuous histopaque density gradient (Sigma, Poole, UK). More details are provided in section 2.2.

3.2.2.4 Cigarette smoke extract preparation

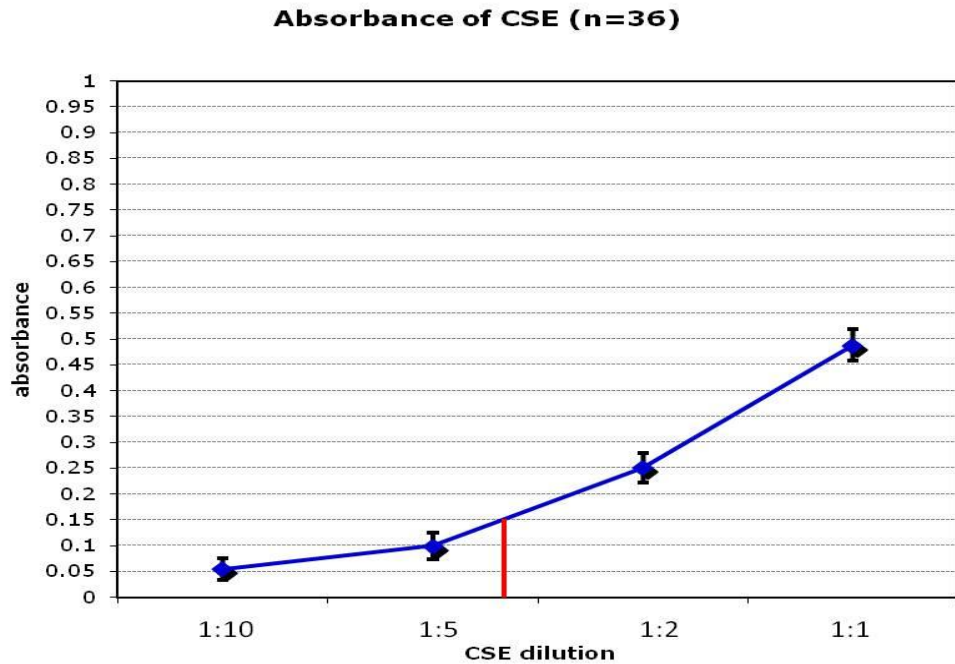


Figure 3.2.1 Absorbance of CSE at different dilutions.

The absorbance of 0.15 is considered as 100% CSE.

CSE were freshly prepared prior to 30 minutes of cell culture for each experiment to avoid breakdown of substances in the extract and evaporation of volatile components. More details are provided in section 2.4. Figure 3.2.1 shows the trend of absorbance according to CSE dilution. Thirty-six (n=36) cigarette were processed in this manner during the whole project.

3.2.2.5 PBMCs culture and treatments

In preliminary experiment to determine the optimal culture time and concentrations of CSE, α -CD3, IL-2 or IL-15 for

Table 3.2.6 Concentrations of stimulus used for activating killer cell.

CSE	0.2%	0.5%	1%	2%	5%
α -CD3	0.1ug/ml	0.2ug/ml	0.5ug/ml	1ug/ml	2ug/ml
IL-2	5ng/ml	10ng/ml	20ng/ml	50ng/ml	100ng/ml
IL-15	1ng/ml	2ng/ml	5ng/ml	10ng/ml	20ng/ml

activation of killer cells, PBMCs isolated from healthy non-smoking participants were resuspended in RPMI-1640 medium supplemented with 10% FCS at a density of 1×10^6 cells/ml and cultured at 200ul/well in 96-well flat bottom tissue culture plates. These cells were either untreated or treated separately with CSE, α -CD3, IL-2 or IL-15 at various concentrations as indicated in Table 3.2.6. Specially, α -CD3 was pre-coated in the plate for 90 minutes at 37°C in 5% CO₂ and washed in cold sterile D-PBS twice to remove the unbound antibody. After incubation for 24 hours, 48 hours and 72 hours, cells were harvested at each time point and washed twice in RPMI-1640 medium for surface marker staining to detect cell activation.

In CSE suppression experiments, PBMCs isolated from HNS, HS, cuS-COPD and exS-COPD were untreated or treated with CSE, α -CD3, IL-2 or IL-15 at each optimized concentrations separately. In order to test the effects of CSE on activated

killer cells, cells stimulated by α -CD3, IL-2 or IL-15 were also co-treated by CSE. For CSE suppression experiment, cells were incubated for 48 hours at 37°C in 5% CO₂ before analysis.

3.2.2.6 Extracellular staining

Fluorescently conjugated antibodies used in this study for extracellular staining were as follows, CD3 (PC7), CD56 (PE), CD8 (ECD) CD25 (FITC) and CD69 (PC5), or KIR (FITC). More details are provided in section 2.6.

3.2.2.7 Flow cytometric analysis

Untreated or treated PBMCs were analysed by flow cytometry. Cells were harvested and fixed in 3% formaldehyde in isotonic azide free solution (Beckman Coulter, Luton, UK) and washed before labelled antibodies were added at the recommended concentration. The cells were incubated in the dark and excess antibody removed. Flow cytometric analysis of antibody labelled cells was performed using an EPICS Altra (Beckman Coulter, Luton, UK). More details are provided in section 2.9.

3.2.2.8 Statistical analysis

The statistical analysis was performed with Prism software, version 4.0c (GraphPad). Differences between the four groups

of participants were tested using the parametric student's *t* test. Confidence interval 99% was used to show two-tailed *p* values. *P* values of less than 0.01 were considered to indicate statistical significance due to the small size of each study group..

3.2.3 Results

3.2.3.1 Optimal concentrations and culture time for killer cell activation

To determine the optimal concentrations of interested stimulus and culture time, PBMCs from healthy non-smokers (*n*=4) were cultured in medium alone or in medium supplemented with either CSE, α -CD3, IL-2 or IL-15 at increasing concentrations (as shown in Table 3.2.6) for 24 hours, 48 hours and 72 hours. Cells are designated activated when they express one or both of the activation markers (i.e., CD69 positive, CD69/CD25 double positive or CD25 positive).

Dose- and time-dependent increases in percentages of killer cell activation were observed during culture with increasing concentrations of α -CD3 (Figure 3.2.2), IL-2 (Figure 3.2.3) and IL-15 (Figure 3.2.4).

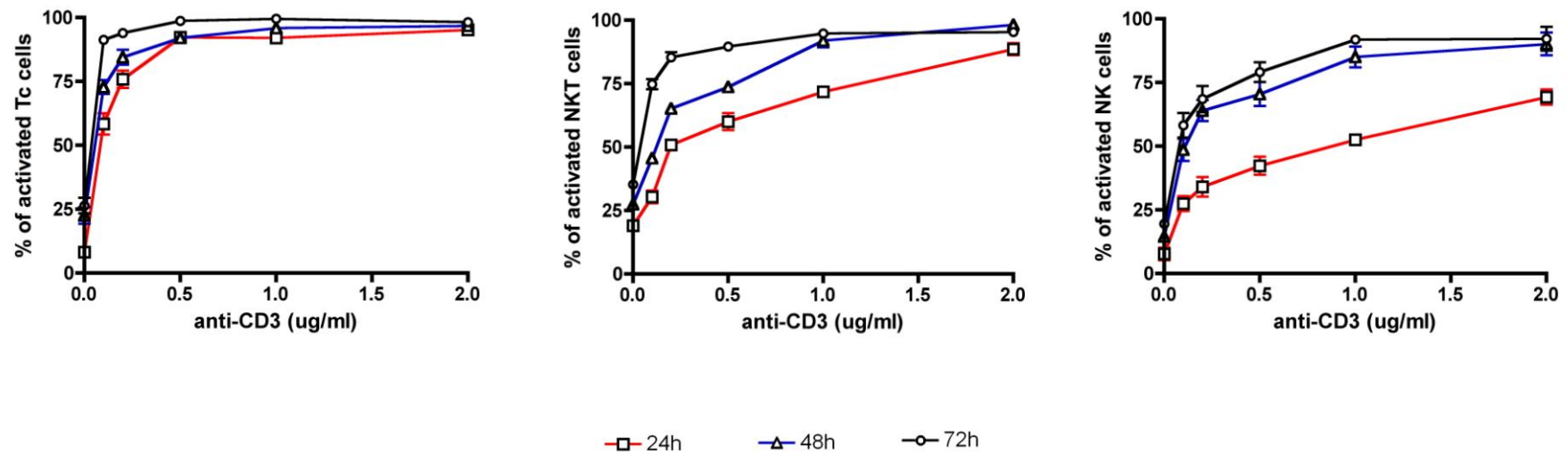


Figure 3.2.2 Activation levels of killer cells by different concentrations of α -CD3. PBMCs were collected from healthy non-smokers (n=4). Cells were untreated or treated by α -CD3 at concentrations of 0.1ug/ml, 0.2ug/ml, 0.5ug/ml, 1ug/ml and 2ug/ml respectively for 24 hours, 48 hours and 72 hours.

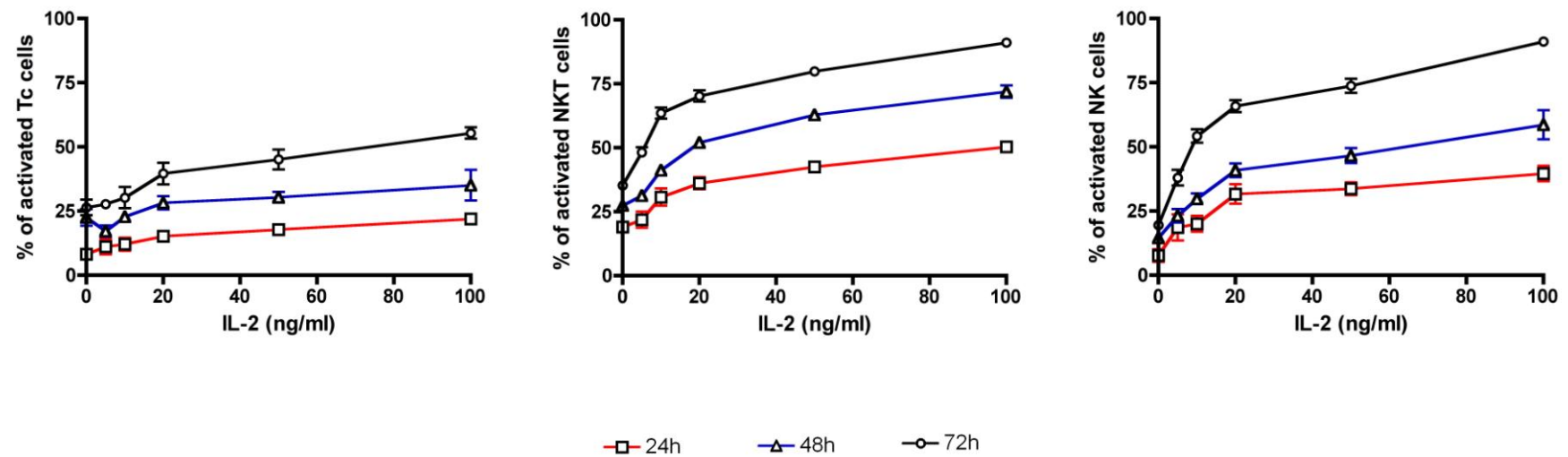


Figure 3.2.3 Activation levels of killer cells by different concentrations of IL-2. PBMCs were collected from healthy non-smokers (n=4). Cells were untreated or treated by IL-2 at concentrations of 5ng/ml, 10ng/ml, 20ng/ml, 50ng/ml and 100ng/ml respectively for 24 hours, 48 hours and 72 hours.

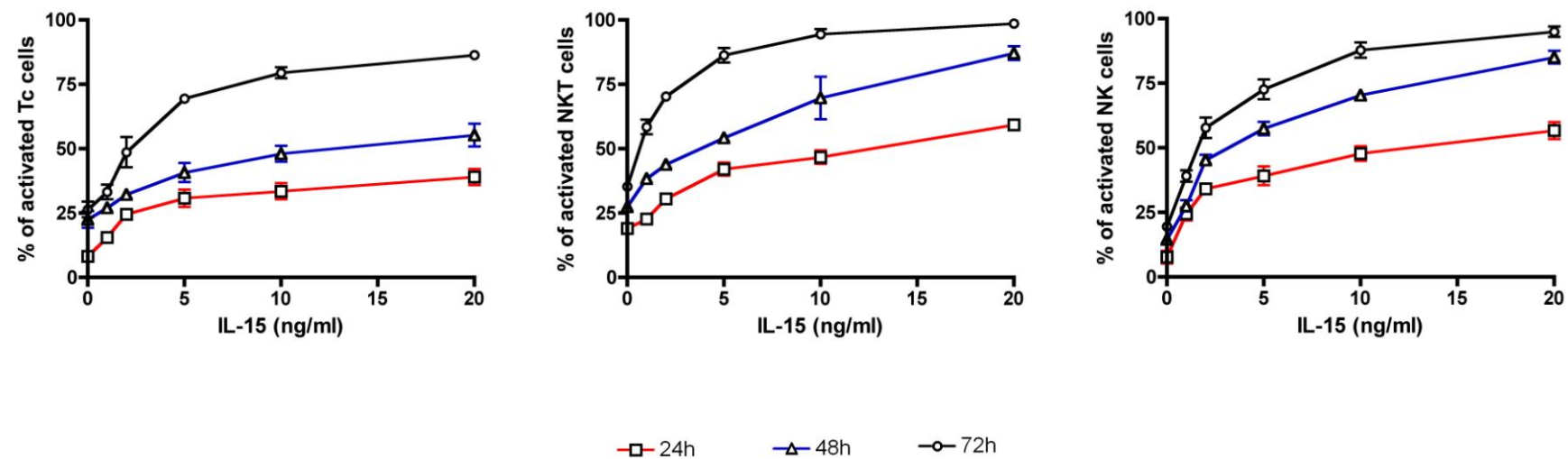


Figure 3.2.4 Activation levels of killer cells by different concentrations of IL-15. PBMCs were collected from healthy non-smokers (n=4). Cells were untreated or treated by IL-15 at concentrations of 1ng/ml, 2ng/ml, 5ng/ml, 10ng/ml and 20ng/ml respectively for 24 hours, 48 hours and 72 hours.

Killer cells of interest including Tc cells, NKT-like cells and NK cells were highly activated when increasing the concentration of α -CD3 to 1ug/ml after incubation for 48 hours (Figure 3.2.2). Activations of killer cells by IL-2 at selected concentrations were not as much activation as α -CD3 treated cells (Figure 3.2.3). However, previous studies have shown that administration of high dose of IL-2 in treatment of a variety of diseases is associated with mortality rates of up to 4% . Therefore, we controlled the maximal concentration of IL-2 at 100ng/ml. IL-15 at 20ng/ml also highly activated these killer cells after 48 hours incubation (Figure 3.2.4).

There were no statistically significant differences on the activation levels of killer cells treated by CSE at different percentage concentrations (all $p > 0.05$) (Figure 3.2.5).

To sum up, to highly activate all killer cells of interest, the optimal concentration of α -CD3, IL-2, IL-15 for detection of cell activation would be chosen at 1ug/ml, 100ng/ml, and 20ng/ml respectively in future experiments. The dilution at 1% of CSE would also be used for following studies. Figure 3.2.6 showed representative dot plots of interested killer cell activation at optimal concentration and dilution after 48 hours incubation.

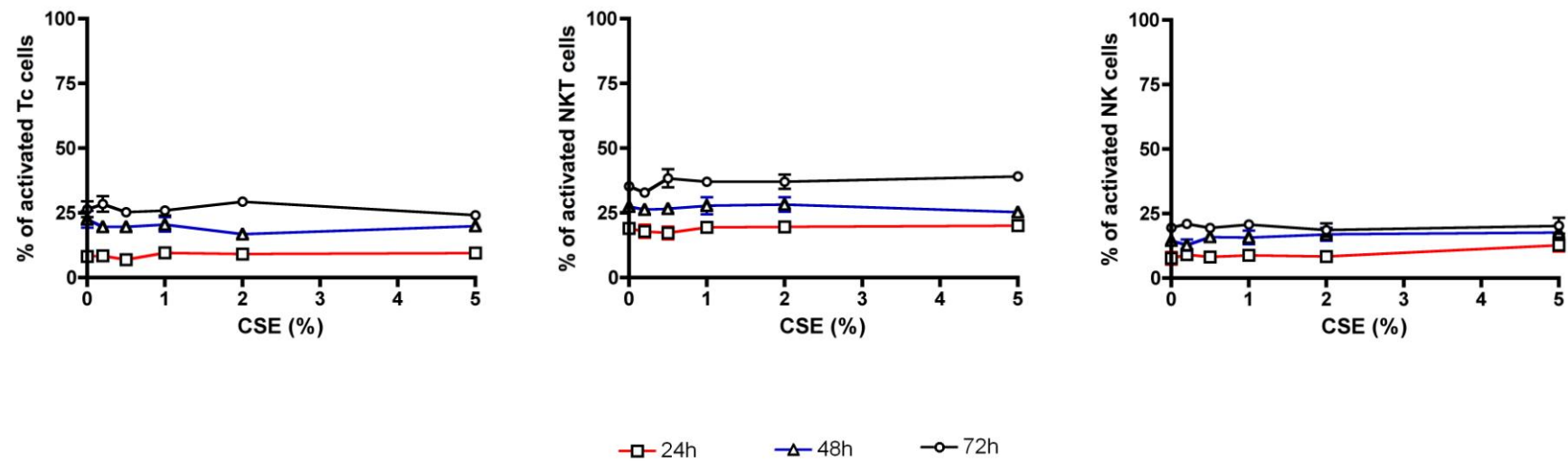


Figure 3.2.5 Activation levels of killer cells by different dilutions of CSE. PBMCs were collected from healthy non-smokers (n=4). Cells were untreated or treated by CSE at dilutions of 0.2%, 0.5%, 1%, 2% and 5% respectively for 24 hours, 48 hours and 72 hours.

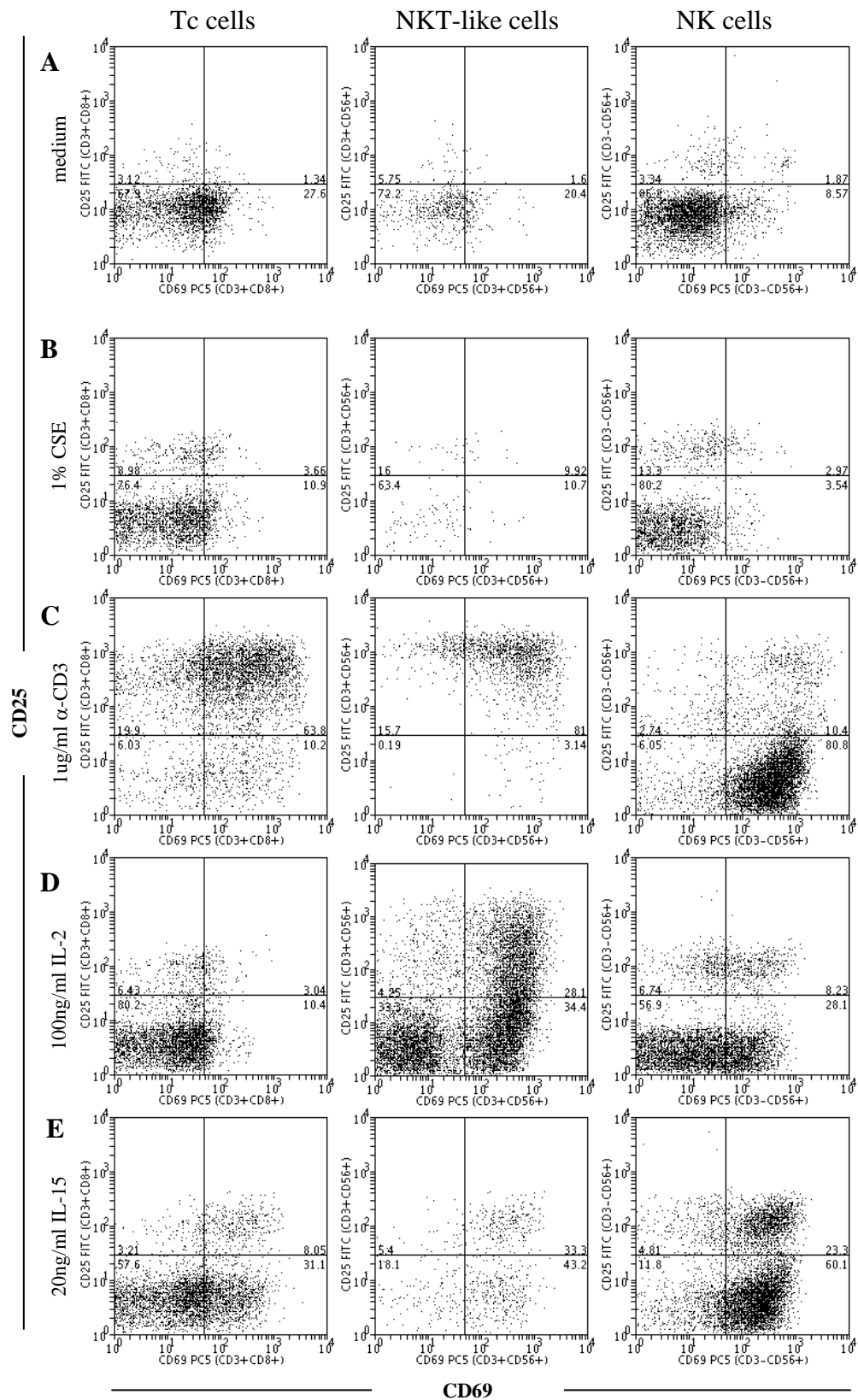
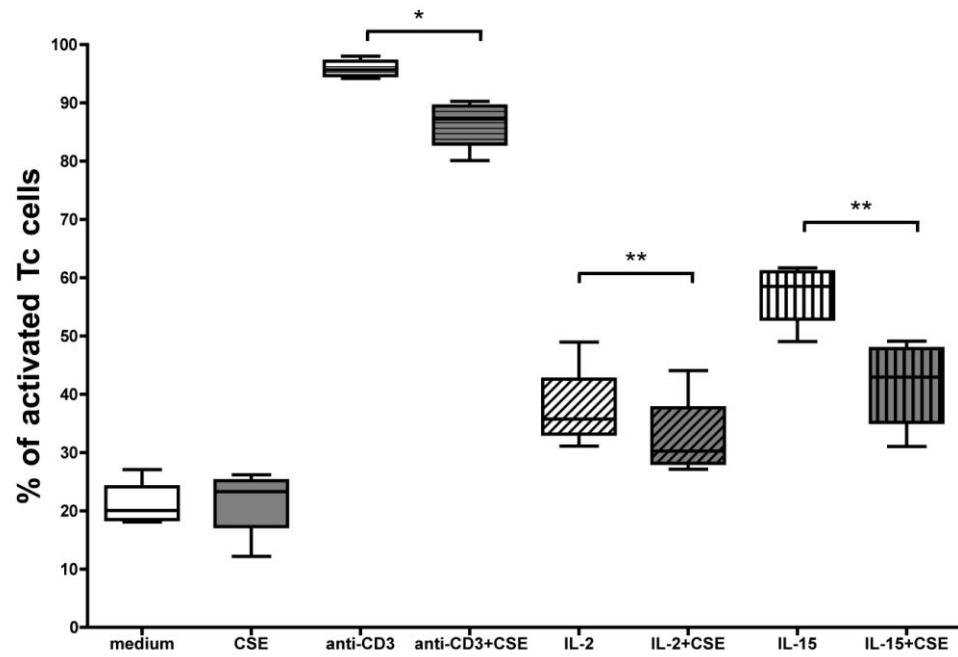


Figure 3.2.6 Representative dot plots for activation of Tc cells, NKT-like cells and NK cells in PBMCs from healthy non-smokers (n=4) by different stimulus at 48 hours. A) medium only; B) 1% CSE; C) 1ug/ml α -CD3; D) 100ng/ml IL-2; and E) 20ng/ml IL-15.

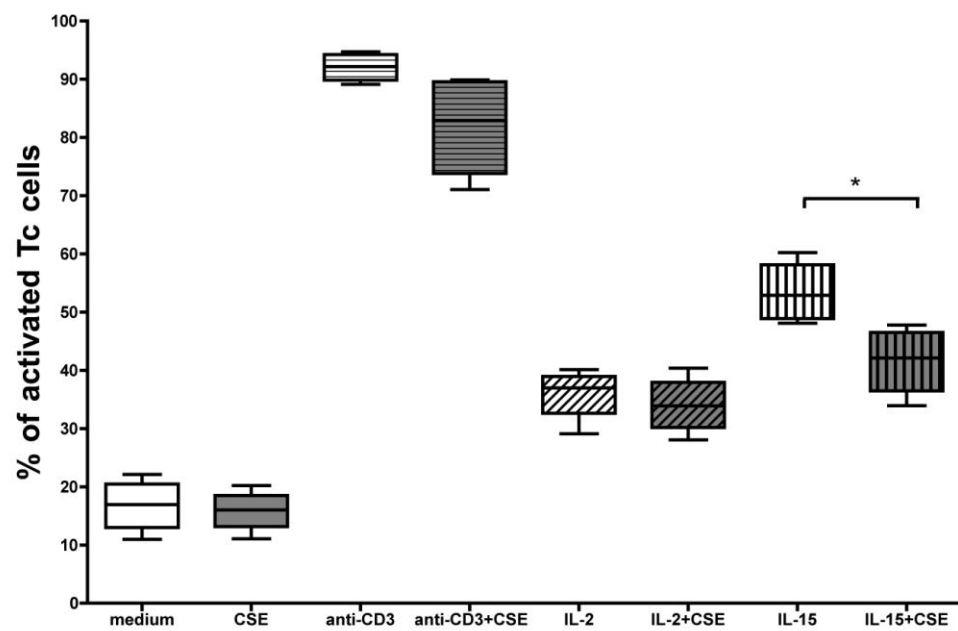
3.2.3.2 CSE suppressed killer cell activation in response to different stimulus

PBMCs collected from the four study groups were either untreated (control) or treated by 1% of CSE alone, or incubated with 1ug/ml α -CD3, 100ng/ml IL-2 or 20ng/ml IL-15 for 48 hours in the absence or presence of 1% of CSE. There was no significant difference on the activation of CD8⁺ T cells (Tc cells) between untreated PBMCs and CSE treated PBMCs in each group (Figure 3.2.7). CSE significantly reduced the activation levels of anti-CD3 treated Tc cells from healthy non-smokers (Figure 3.2.7.A, $p < 0.01$) and current smokers with COPD (Figure 3.2.7.C, $p < 0.005$). Also, IL-2 induced Tc cell activation was dramatically decreased when CSE was added in healthy non-smokers (Figure 3.2.7.A, $p < 0.005$), current smokers with COPD (Figure 3.2.7.C, $p < 0.001$) and ex-smokers with COPD (Figure 3.2.7.D, $p < 0.001$). Furthermore, CSE markedly attenuated IL-15 stimulated T cell activation in all interested groups (p value was shown in Table 3.2.7).

A. Healthy non-smokers (HNS)

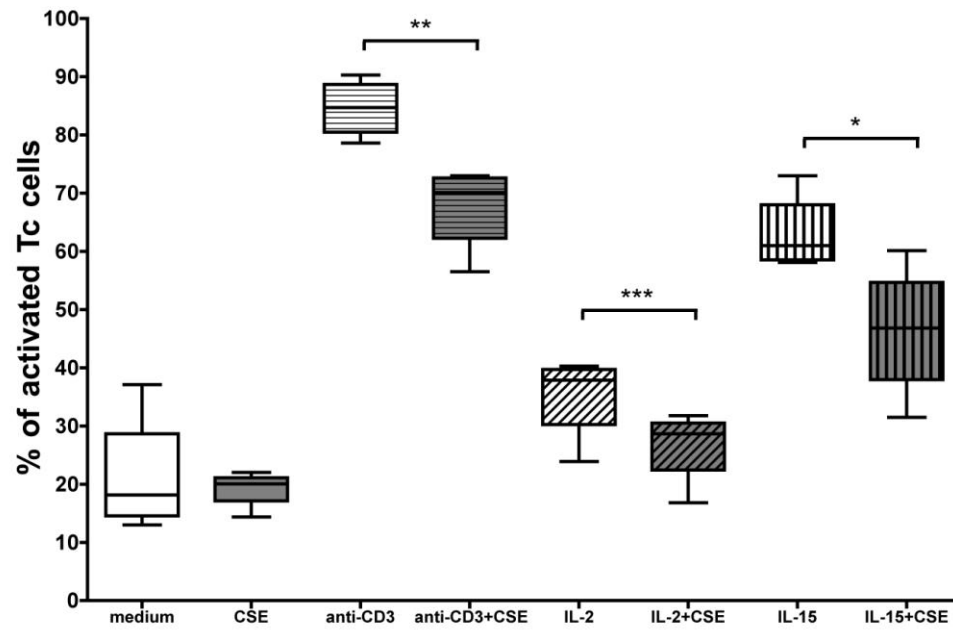


B. Current Healthy smokers (HS)



CSE (1%); anti-CD3 (1ug/ml)
IL-2 (100ng/ml); IL-15 (20ng/ml)

C. current smokers with COPD (cuS-COPD)



D. ex-smokers with COPD (exS-COPD)

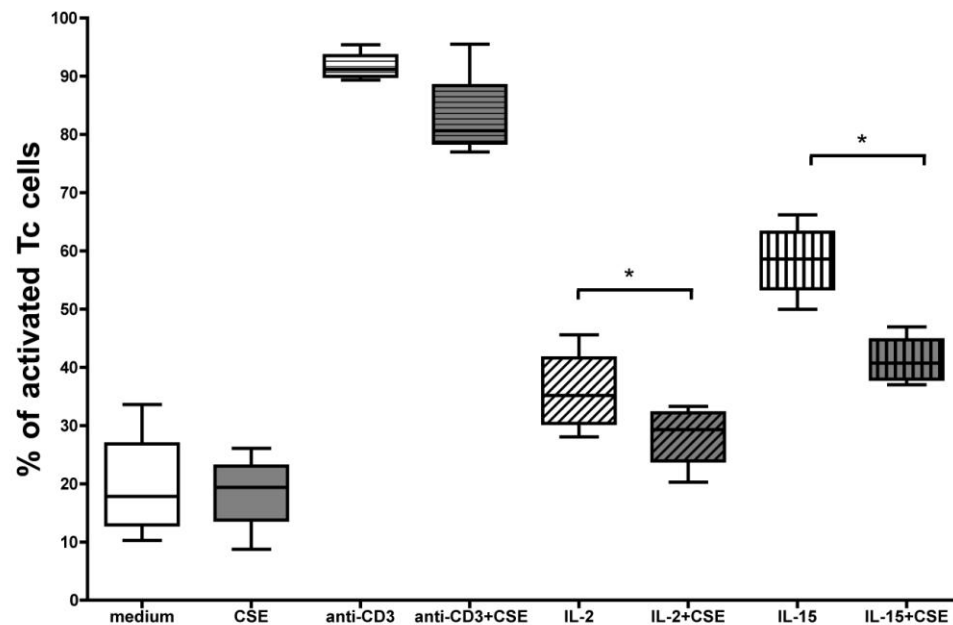


Figure 3.2.7 CSE effects on activation levels of Tc cells in four groups. PMBCs were collected from four groups including A) HNS (n=4), B) HS (n=4), C) cuS-COPD (n=4), and D) exS-COPD (n=4). Cells were untreated or treated by different reagents at designed concentrations and incubated 48 hours. *, $p < 0.01$; **, $p < 0.005$; ***, $p < 0.001$.

Table 3.2.7 Statistical difference of before and after CSE treated cells

		Medium vs CSE	anti-CD3 vs anti-CD3+CSE	IL-2 vs IL-2+CSE	IL-15 vs IL-15+CSE
Tc cells	HNS	p>0.01	p<0.01	p<0.005	p<0.005
	HS	p>0.01	p>0.01	p>0.01	p<0.01
	cuS-COPD	p>0.01	p<0.005	p<0.001	p<0.01
	exS-COPD	p>0.01	p>0.01	p<0.01	p<0.01
NKT-like cells	HNS	p>0.01	p<0.005	p<0.01	p<0.01
	HS	p>0.01	p>0.01	p>0.01	p<0.005
	cuS-COPD	p>0.01	p<0.005	p<0.005	p<0.01
	exS-COPD	p>0.01	p<0.005	p<0.01	p<0.005
NK cells	HNS	p>0.01	p<0.001	p<0.01	p<0.005
	HS	p>0.01	p<0.005	p<0.005	p<0.001
	cuS-COPD	p>0.01	p<0.005	p<0.005	p<0.001
	exS-COPD	p>0.01	p<0.005	p<0.005	p<0.001

- HNS, healthy non-smokers; HS, current healthy smokers; cuS-COPD, current smokers with COPD; exS-COPD, ex-smokers with COPD.
- CSE showed significant effects on all cell types of interest upon IL-15 stimulation in all groups.

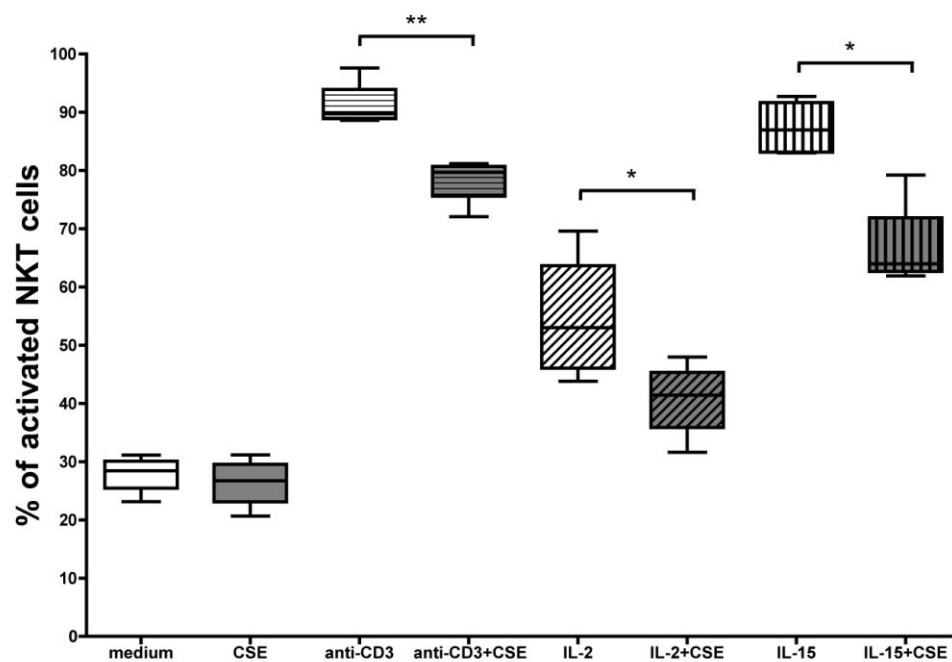
CSE effects on activated NKT cells were also investigated. There was no significant difference on the activation levels of NKT-like cells treated with or without CSE in each group (Figure 3.2.8). CSE significantly reduced the activation levels of α -CD3 treated Tc cells from healthy non-smokers (Figure 3.2.8.A, p<0.005), current smokers with COPD (Figure 3.2.8.C, p<0.005) and ex-smokers with COPD (Figure 3.2.8.D, p<0.005). IL-2 induced NKT-like cell activation was dramatically decreased when CSE was added in healthy non-smokers (Figure 3.2.8.A, p<0.01), current smokers with COPD

(Figure 3.2.8.C, $p < 0.005$) and ex-smokers with COPD (Figure 3.2.8.D, $p < 0.01$). Furthermore, CSE markedly attenuated IL-15 stimulated NKT-like cell activation in all groups (p value was shown in Table 3.2.7).

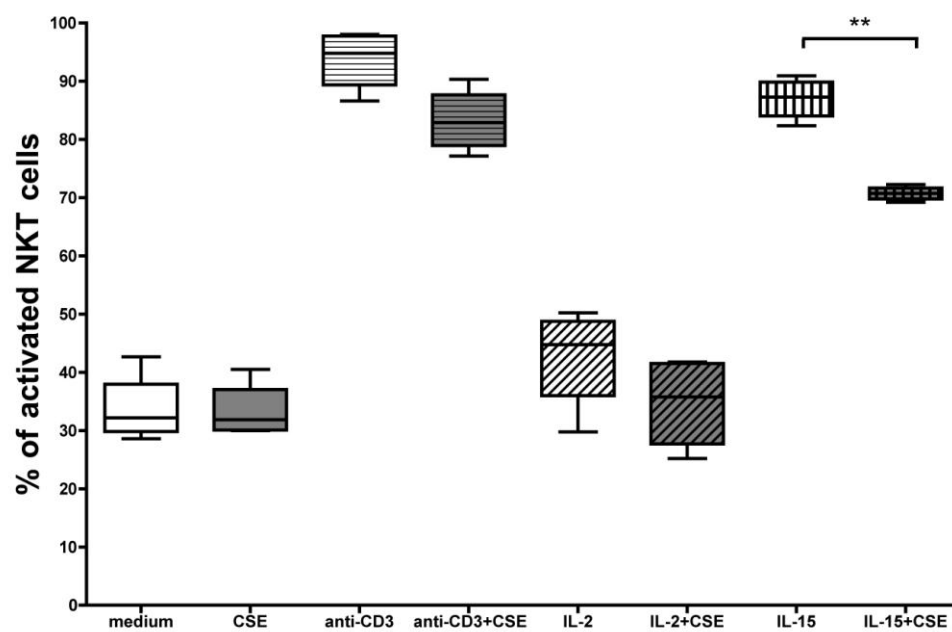
When examining activated levels on NK cells, untreated cells compared to CSE alone had no markedly effects on NK cell activation in each patient group (Figure 3.2.9). Activation levels of NK cell in response to α -CD3, IL-2 or IL-15 were all significantly decreased in the presence of CSE (p value was shown in Table 3.2.7).

Absolute activation levels of killer cells in response to each stimulation were also compared between groups. There were no statistical differences detected (Figures are shown in Appendix).

A. Healthy non-smokers (HNS)

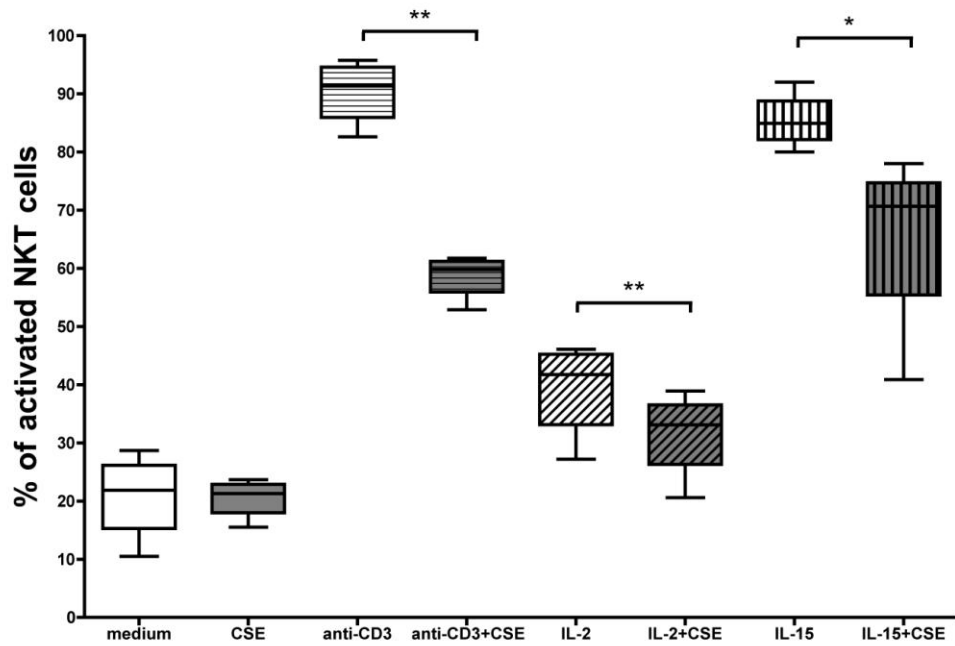


B. Current healthy smokers (HS)



CSE (1%); anti-CD3 (1ug/ml)
IL-2 (100ng/ml); IL-15 (20ng/ml)

C. current smokers with COPD (cuS-COPD)



D. ex-smokers with COPD (exS-COPD)

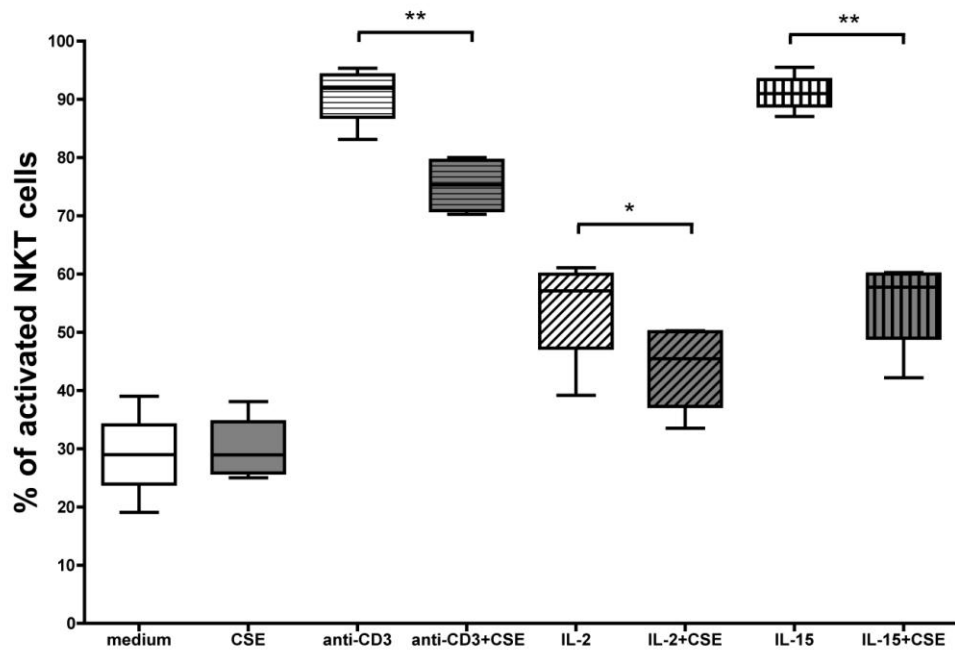
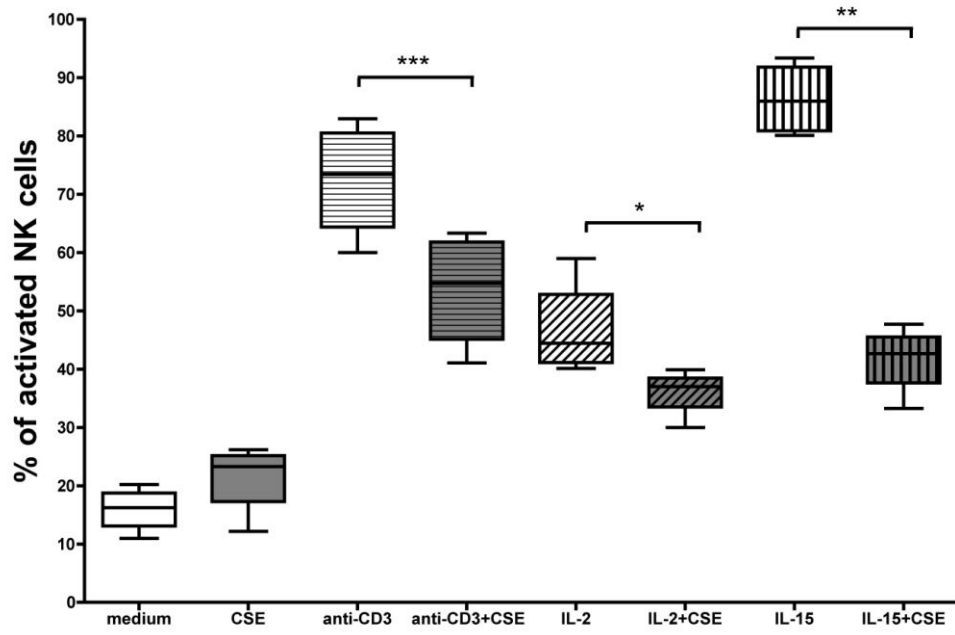
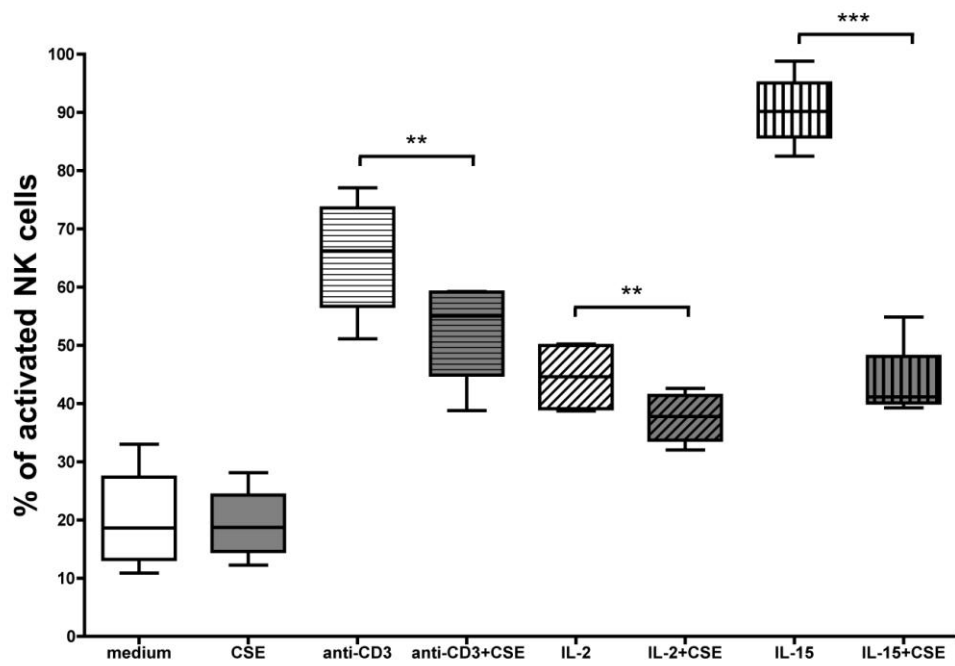


Figure 3.2.8 CSE effects on activation levels of NKT cells in four groups. PMBCs were collected from four groups including A) HNS (n=4), B) HS (n=4), C) cuS-COPD (n=4), and D) exS-COPD (n=4). Cells were untreated or treated by different reagents at designed concentrations and incubated 48 hours. *, $p < 0.01$; **, $p < 0.005$; ***, $p < 0.001$.

A. Healthy non-smokers (HNS)

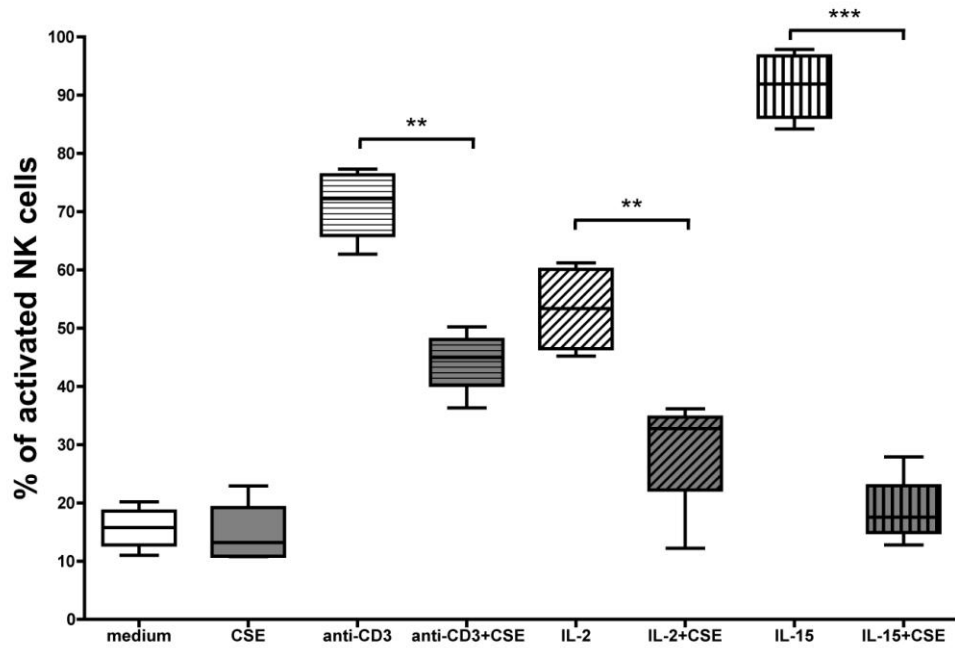


B. Current healthy smokers (HS)



CSE (1%); anti-CD3 (1ug/ml)
IL-2 (100ng/ml); IL-15 (20ng/ml)

C. current smokers with COPD (cuS-COPD)



D. ex-smokers with COPD (exS-COPD)

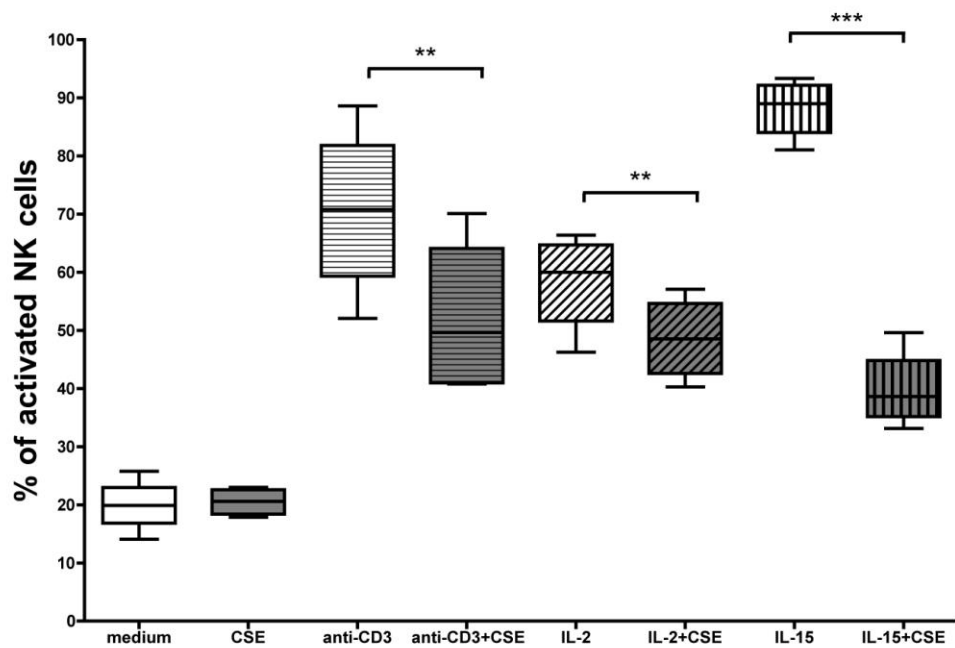


Figure 3.2.9 CSE effects on activation levels of NK cells in four groups. PMBCs were collected from four groups including A) HNS (n=4), B) HS (n=4), C) cuS-COPD (n=4), and D) exS-COPD (n=4). Cells were untreated or treated by different reagent at designed concentrations and incubated 48 hours. *, $P < 0.01$; **, $p < 0.005$; ***, $p < 0.001$.

3.2.3.3 Absolute suppression levels induced by CSE on killer cell activation in response to different stimulation

Since CSE dramatically suppressed most activation levels of Tc cells, NKT-like cells and NK cells upon different stimulation in each group, we further compared the absolute suppression levels induced by CSE on cell activation between four groups. The calculation formula of absolute suppression levels was "Absolute suppression level = the percentage of activated cells of interest up on different stimulation - the percentage of these cells upon CSE treatment". There were no significant differences on the absolute levels of CSE induced suppression in α -CD3 treated Tc cell activation between four groups (Figure 3.2.10.A, all $p > 0.01$). Absolute suppression level of CSE treated NKT cells upon α -CD3 stimulation was markedly higher in current smokers with COPD compared to healthy non-smokers ($p < 0.005$), current healthy smokers ($p < 0.01$) and ex-smokers with COPD ($p < 0.005$) (Figure 3.2.10.B). Furthermore, CSE inducing absolute suppression level on NK cell activation upon α -CD3 stimulation was markedly higher in current smokers with COPD compared to healthy non-smokers ($p < 0.01$) and current healthy smokers ($p < 0.005$) (Figure 3.2.10.C). The absolute suppression level on CSE treated NK cells in response to α -CD3 was also significantly

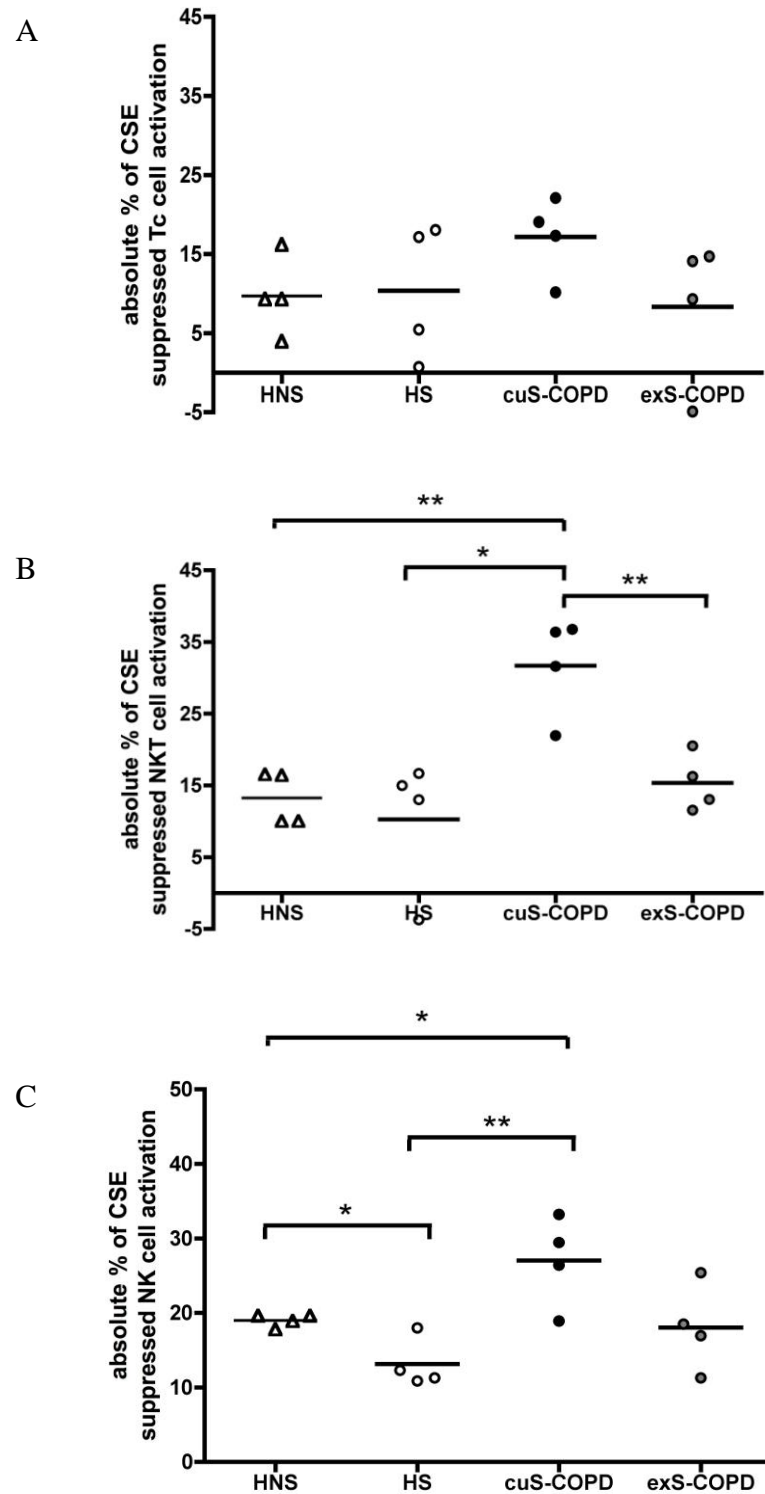


Figure 3.2.10 Absolute suppression levels induced by CSE on α -CD3 treated killer cell activation in four groups. A) Tc cells; B) NKT cells; C) NK cells. HNS (healthy non-smokers, n=4); HS (current healthy smokers, n=4); C) cuS-COPD (current smokers with COPD, n=4); exS-COPD (ex-smokers with COPD, n=4). *, $P < 0.01$; **, $p < 0.005$; ***, $p < 0.001$.

higher in healthy non-smokers ($p<0.005$) in contrast to current healthy smokers (Figure 3.2.10.C).

In IL-2 treated Tc cells, absolute suppression levels induced by CSE were significantly higher in current smokers with COPD compared to healthy non-smokers ($p<0.005$) and current healthy smokers ($p<0.01$) (Figure 3.2.11.A). Also, absolute suppression level on CSE treated NK cell activation was remarkably higher in current smokers with COPD compared to healthy non-smokers ($p<0.01$), current healthy smokers ($p<0.005$) and ex-smokers with COPD ($p<0.005$) (Figure 3.2.11.C). There were no significant differences on the absolute levels of CSE inducing suppression on Tc cell activation between four groups (Figure 3.2.11.B, all $p>0.01$).

CSE did not show any statistical differences on the absolute suppression levels of IL-15 stimulated Tc cells between groups (Figure 3.2.12.A). NKT cells in response to IL-15 was dramatically suppressed when CSE was added in ex-smokers with COPD compared to healthy non-smokers ($p<0.01$) and current healthy smokers ($p<0.001$) (Figure 3.2.12.B). Moreover, the absolute suppression level induced by CSE on IL-15 treated NK cells was significantly higher in current smokers with COPD compared to other three groups including

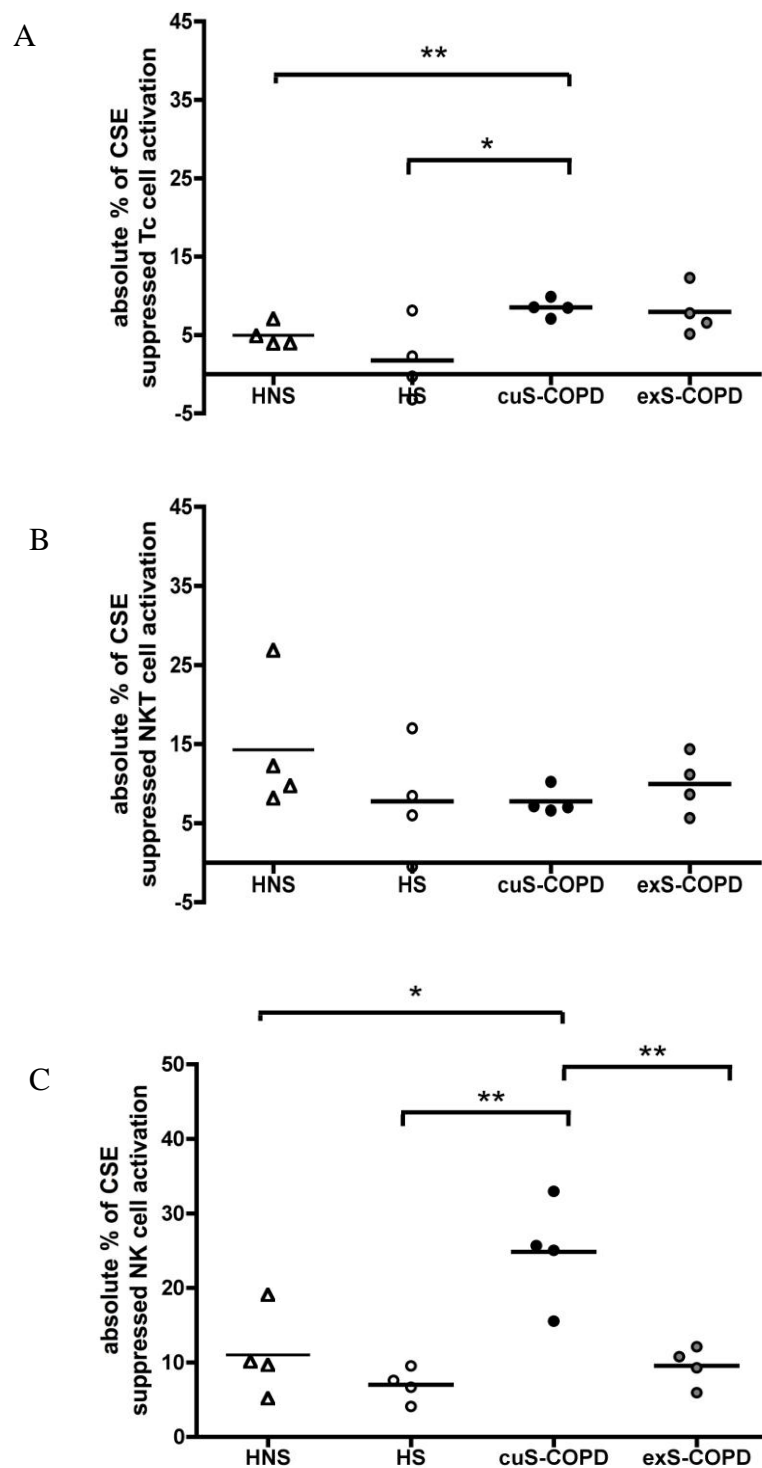


Figure 3.2.11 Absolute suppression levels induced by CSE on IL-2 treated killer cell activation in four groups. A) Tc cells; B) NKT cells; C) NK cells. HNS (healthy non-smokers, n=4); HS (current healthy smokers, n=4); C) cuS-COPD (current smokers with COPD, n=4); exS-COPD (ex-smokers with COPD, n=4). *, $P < 0.01$; **, $p < 0.005$; ***, $p < 0.001$.

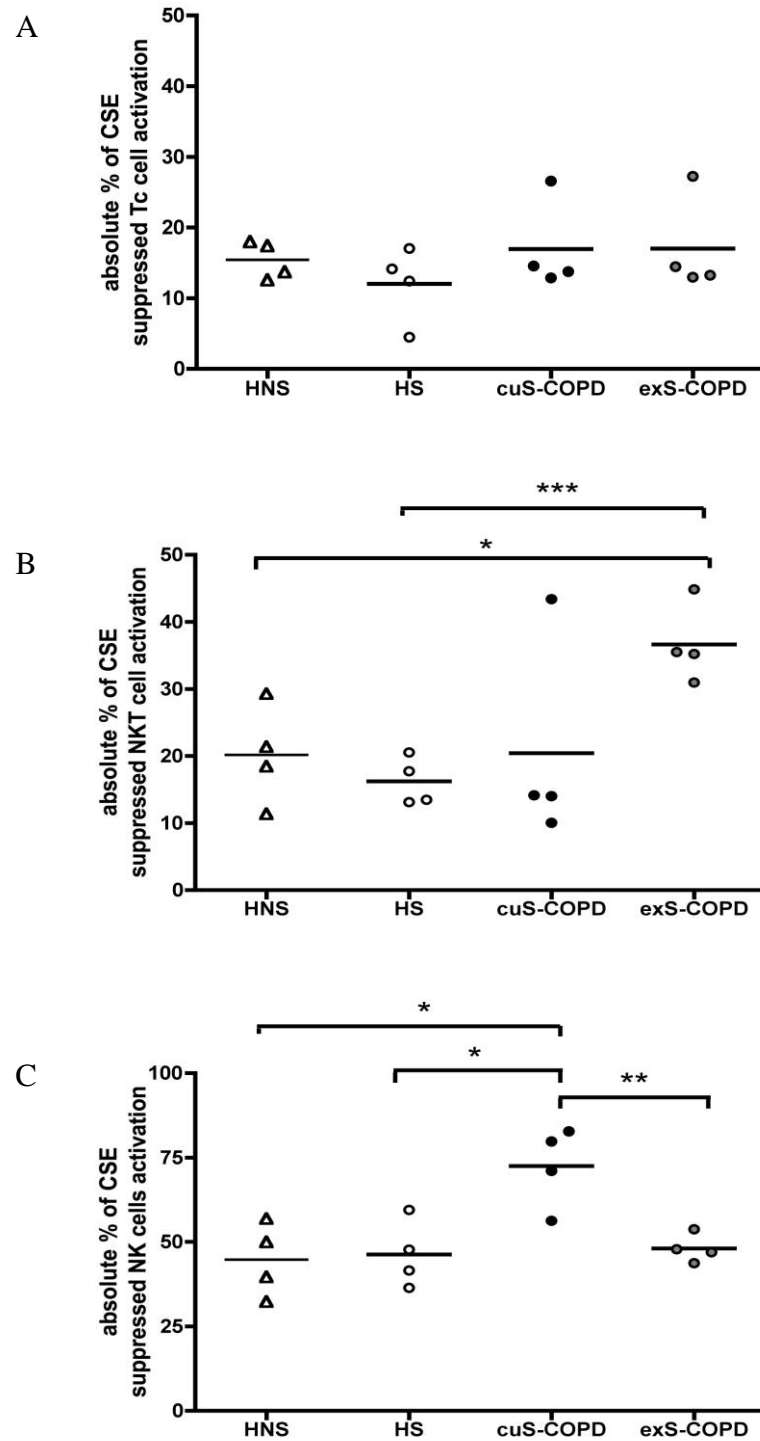


Figure 3.2.12 Absolute suppression levels induced by CSE on IL-15 treated killer cell activation in four groups. A) Tc cells; B) NKT cells; C) NK cells. HNS (healthy non-smokers, n=4); HS (current healthy smokers, n=4); C) cuS-COPD (current smokers with COPD, n=4); exS-COPD (ex-smokers with COPD, n=4). *, $P < 0.01$; **, $p < 0.005$; ***, $p < 0.001$.

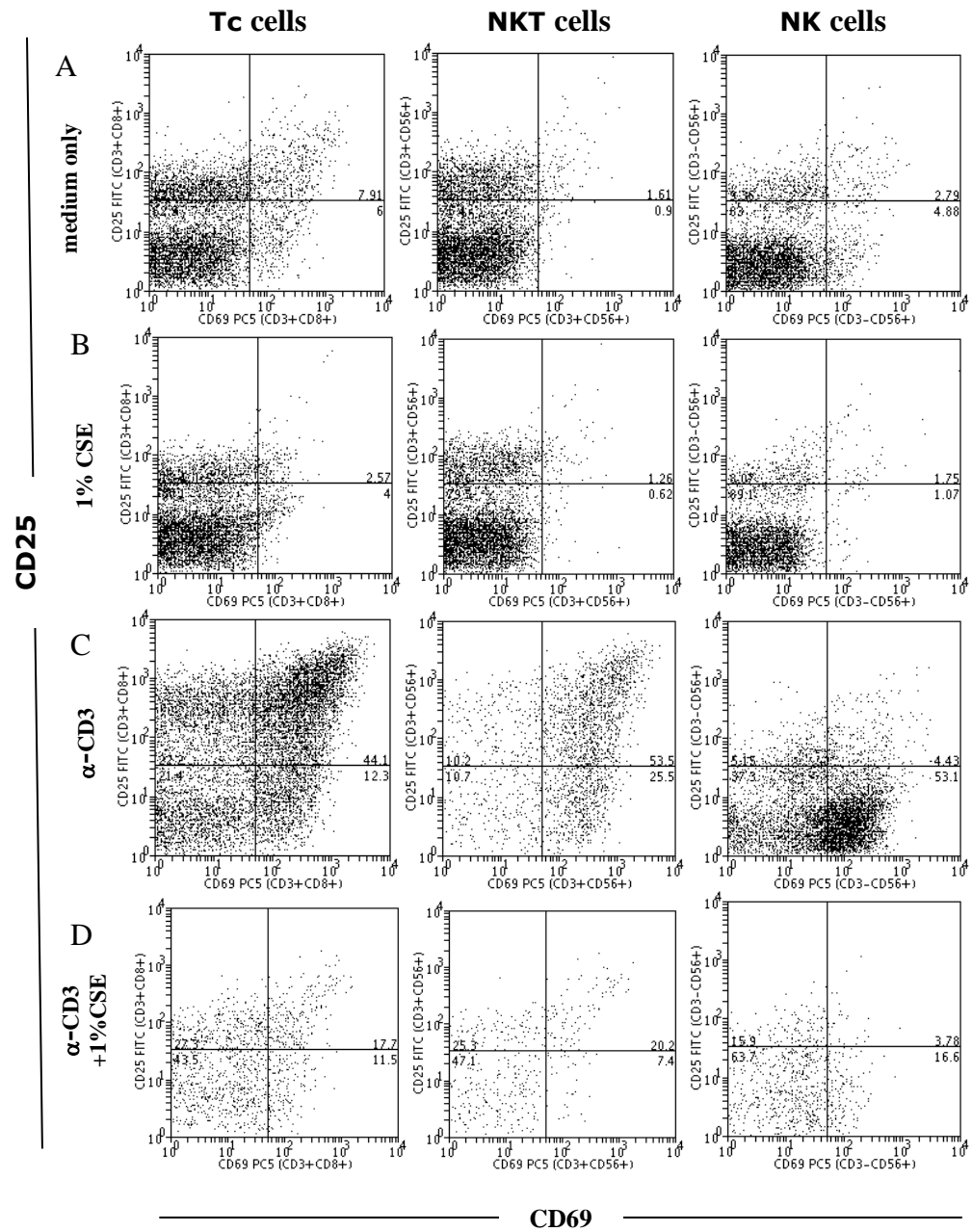
healthy non-smokers ($p < 0.01$), current healthy smokers ($p < 0.01$) and ex-smokers with COPD ($p < 0.005$) (Figure 3.2.12.C).

Figure 3.2.13 showed the representative dot plots of CSE suppressed cell activation upon different activators from a current smoker with COPD. Figure 1, Figure 2 and Figure 3 from Appendix showed the comparison of absolute activation levels of cells of interest upon different stimulation in four groups. Table 3.2.8 showed the statistical differences of absolute suppression induced by CSE on different cell activation between groups.

3.2.4 Discussion

Cigarette smoking is still the major risk factor in the development of COPD. Accumulating evidence has suggested that chronic inhalation of cigarette smoke alters a wide range of immunologic function, including both innate and adaptive immune responses (Honda et al., 1996, Mio et al., 1997, Mohan, 2002b, Subramaniam et al., 1996, Wyatt et al., 1999). The main focus of this section was to investigate the impact of CSE on α -CD3, IL-2 or IL-15 activated killer cells including Tc cells, NKT-like cells and NK cells, and further compared their activation levels in four groups, namely, healthy non-smokers,

current healthy smokers, current smokers with COPD and ex-smokers with COPD.



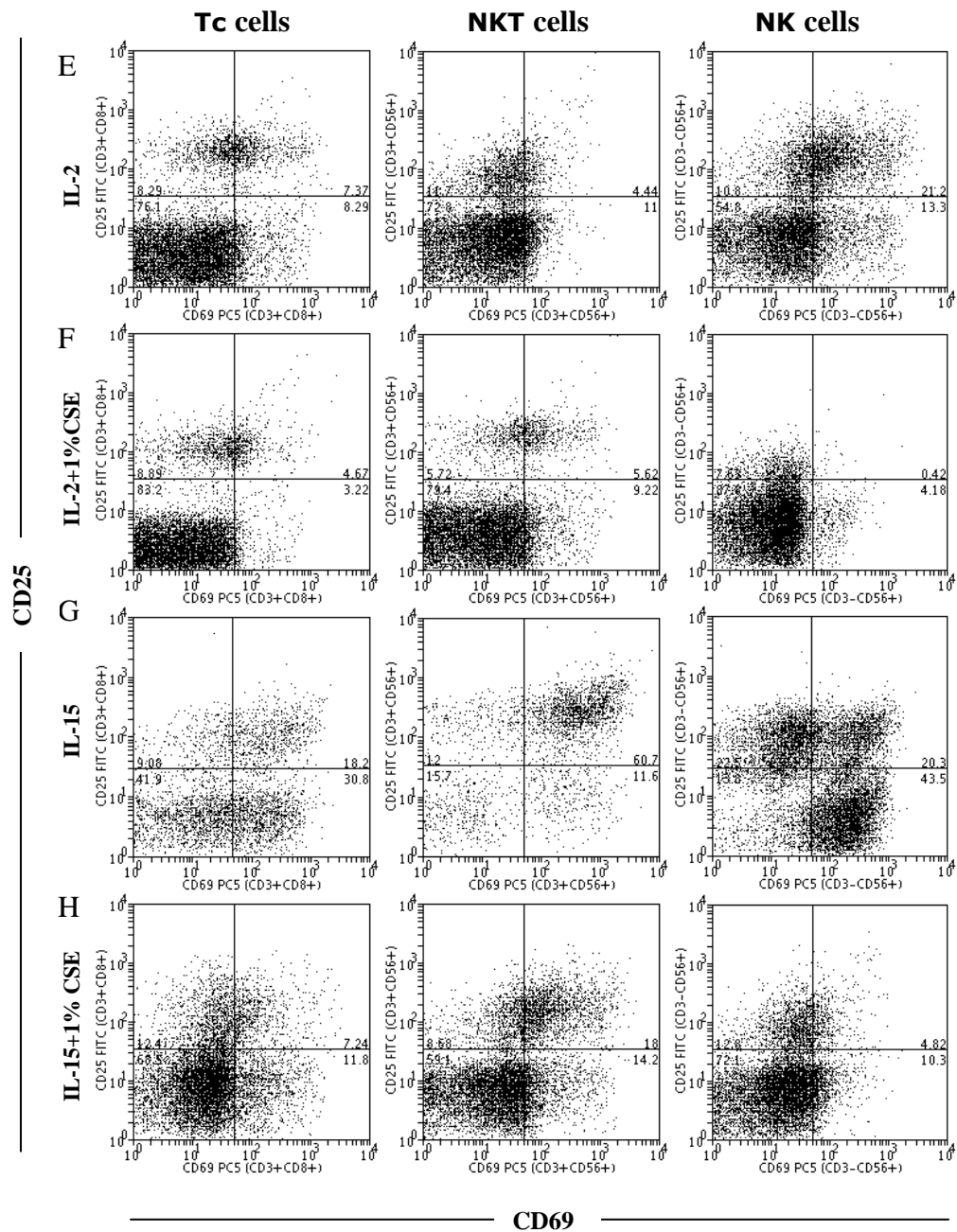


Figure 3.2.13 Representative dot plots of the activation of interested cells from a current smoker with COPD. PBMCs were stimulated by either A) meidium only; B) 1% CSE only; C) α -CD3; D) α -CD3+1% CSE; E) IL-2; F) IL-2+1% CSE; G) IL-15; or H) IL-15+1% CSE; and incubated for 48 hours after stimulation. α -CD3 significantly activated both Tc cells and NKT cells compared to NK cells. Whereas, IL-2 and IL-15 showed remarkable effects on NK cell activation.

Table 3.2.8 Absolute suppression induced by CSE on activation of killer cells upon different stimulation.

	HNS vs HS	HNS vs cuS-COPD	HNS vs exS-COPD	HS vs cuS-COPD	HS vs exS-COPD	cuS-copd vs exS-COPD
α-CD3 treated cells						
Tc	p>0.01	p>0.01	p>0.01	p>0.01	p>0.01	p>0.01
NKT	p>0.01	p<0.005	p>0.01	p<0.01	p>0.01	p<0.005
NK	p<0.01	p<0.01	p>0.01	p<0.005	p>0.01	p>0.01
IL-2 treated cells						
Tc	p>0.01	p<0.005	p>0.01	p<0.01	p>0.01	p>0.01
NKT	p>0.01	p>0.01	p>0.01	p>0.01	p>0.01	p>0.01
NK	p>0.01	p<0.01	p>0.01	p<0.005	p>0.01	p<0.005
IL-15 treated cells						
Tc	p>0.01	p>0.01	p>0.01	p>0.01	p>0.01	p>0.01
NKT	p>0.01	p>0.01	p<0.005	p>0.01	p<0.001	p>0.01
NK	p>0.01	p<0.005	p>0.01	p<0.01	p>0.01	p<0.005

HNS, healthy non-smokers; HS, current healthy smokers; cuS-COPD, current smokers with COPD; exS-COPD, ex-smokers with COPD.

Firstly, to achieve optimal activation of all cells of interest, we set up a preliminary protocol to decide the optimal concentrations of each activator and incubation time for cell culture. Tc cells, NKT-like cells and NK cells were all highly activated when the concentration of α -CD3 increased to 1ug/ml. CD3 is a cell surface marker for T cells including Tc cells, Th cells, Treg cell, γ/δ T cells and NKT cells (Call and Wucherpfennig, 2004). However, in preliminary experiments, dose-dependent increases in the activation levels of NK cells were also observed, although NK cells do not express CD3/TCR complex on their cell surface. To our knowledge, this

could be easily explained. α -CD3 has the capacity of activating CD4⁺ T cells, Tc cells and NKT cells that further release immune mediators such as IL-2 or IL-15 (Smyth et al., 2005, Thompson et al., 1989). Furthermore, previous studies have shown that CSE also has effects on DCs and macrophages that also secrete high amount of IL-15 (Carson et al., 1995, Jonuleit et al., 1997). Both IL-2 and IL-15 are good sources for NK cell activation (Waldmann, 2006).

Activation of killer cells induced by IL-2 is not as great as IL-15 treated cells in healthy non-smokers. One of the reasons is the biological difference on features between IL-2 and IL-15. Although IL-2 and IL-15 share two receptors therefore have certain common features in their functions, they also provide distinct contributions on cell activation and apoptosis (Thomas, 2008, Waldmann and Tagaya, 1999). IL-2R α is only expressed by pre B cells, effector T cells and DCs, whereas IL-15R α is widely expressed on many cell types except regulatory T cells. Furthermore, IL-15 increases the proliferation of the CD8⁺ T cells and reduces their death by apoptosis (Ku et al., 2000, Zhang et al., 1998), whereas IL-2 plays a critical role in Fas-mediated activation-induced cell death (Lenardo, 1996). Kanegane and Tosato has shown that in PBMCs from healthy volunteers, IL-2 and IL-15 can both activate memory CD4⁺ T

cells, memory CD8⁺ T cells and naive CD8⁺ T cells, but not naive CD4⁺ T cells (Kanegane and Tosato, 1996). In addition, IL-2 does not act on resting NKT cells or NK cells that without IL-2R α expression (Waldmann, 2002). We controlled the maximal concentration of IL-2 at 100ng/ml because of the side effects caused by high dose IL-2 administration as previously mentioned.

In our study, activation levels of killer cells were not significantly different in the absence or presence of CSE at increasing dilutions in healthy volunteers, without other stimulations. These findings were in agreement with other groups who detected the CSE effects on DCs or NK cells from healthy volunteers (Mian et al., 2009, Mortaz et al., 2009). Although Stringer *et al* showed that CSE impaired neutrophil caspase-3-like activity in a dilution-dependent manner (CSE dilution 2%-16%), they were concerned about different cell types and the incubation time was only for 4 hours (Stringer et al., 2007). According to previous studies and our detection, we selected 1% of CSE for further experiments because it did not cause any toxic effects as revealed by MMT assay (Mian et al., 2008, Mian et al., 2009, Stringer et al., 2007).

The effects of CSE on killer cells were explored not only in healthy volunteers but also COPD patients. Although CSE alone did not show any impacts on the killer cells activation in all interested groups, we here for the first time showed that CSE significantly reduced the activation levels of Tc cells, NKT cells and NK cells in response to α -CD3, IL-2 or IL-15 in different groups (Table 3.2.7). Of note, three representative results should be pointed out. Firstly, CSE markedly suppressed activation levels of IL-15 treated all cell types in all groups. Secondly, in contrast to Tc cells and NKT cells, NK cells were the only cell type which activation levels upon different stimulation were all significantly decreased when CSE was added in all participant groups. Thirdly, current smokers with COPD were the only participant group that showed CSE had statistically significant effects on all killer cell type activation with the different stimulations. These findings were supported by different studies. Mian and colleague have shown that in PBMCs from healthy non-smokers, the expression of CD69, a key activation marker for NK cells, was significantly inhibited when PBMCs are treated by cigarette smoke after poly I:C activation (Mian et al., 2009). Also, the cigarette smoke markedly reduced NK cell cytolytic potential to kill K562 cancer cells which was found to be IL-15 mediated. Recently, Hogan *et al* demonstrated that CSE significantly

reduced the cytokine production and the ability to kill target cells of α -GalCel activated invariant NKT cells in PBMCs from healthy non-smoking participants. Meanwhile, the activation marker of CD107, but not CD69, was decreased (Hogan et al., 2011). Currently there are no previous studies to show the CSE effects on anti-CD3 activated killer cells.

Since CSE showed suppression effects on killer cells at different levels in each group, we further compared suppression levels in each cell type between healthy volunteers and patient groups. Due to the diverse background of each sample, we used absolute suppression levels to find out whether they have statistical differences. There were no significant differences of CSE suppression on Tc cells in response to α -CD3 or IL-15 between all groups. Interestingly, the absolute suppression levels induced by CSE on IL-2- or IL-15-treated NK cell activation were significantly higher in current smokers with COPD compared to other three groups, namely, healthy non-smokers, current healthy smokers and ex-smokers with COPD. In α -CD3 stimulated NKT cells, their activation levels were also dramatically suppressed when CSE was added in current smokers with COPD compared to healthy participants and ex-smokers with COPD. However, CSE remarkably reduced the activation levels of NKT cells upon IL-

15 activation in ex-smokers with COPD, but not current smokers with COPD, compared to healthy non-smokers and current healthy smokers.

Chronic exposure to cigarette smoke leads to not only cellular death and damage in the airways, but also failure to activate critical intracellular signalling mechanisms that initiate protective host defence. The consequence is dysfunction of various cell types with accompanying predisposition to respiratory infection (Feldman and Anderson, 2013). Although small sample size was studied, our findings showed that in different conditions, CSE indeed suppressed killer cell activation in either healthy groups or COPD patients, or all of the samples. We believe in future, more reliable and consistent findings will be revealed when the sample size is enlarged.

It is important to note that cigarette smoke comprises a complex mixture of more than 4000 chemicals, such as nicotine, carbon monoxide, and tar (Hoffmann and Wynder, 1986, Stedman, 1968). Their immune-modulation effects on immune cells are distinct and also dose-dependent (Holt et al., 1976). To determine the specific component of cigarette

smoke that regulates killer cell activation upon IL-15 stimulation is beyond the scope of our studies.

In conclusion, we here for the first time demonstrate that absolute suppression levels induced by CSE on IL-2- or IL-15-mediated NK cell activation were significantly higher in current smokers with COPD patients compared to healthy non-smokers, current healthy smokers and ex-smokers with COPD. These findings provide further insights on understanding the cellular mechanisms of COPD and may also explain why smokers are more susceptible to lung infections.

3.3 Activation and cytotoxic molecule expression of CD56⁺ cells from peripheral blood samples

3.3.1 Introduction

The previous chapter demonstrated that CSE significantly suppressed NK cell activation in IL-15 treated PBMCs, especially in COPD patients. Activation levels of NKT-like cells upon IL-15 stimulation were also markedly decreased in ex-smokers with COPD. Collectively, these findings suggested that CSE had impact on IL-15 mediated CD56⁺ cell activation.

One of the major killing mechanisms of cytotoxic cells is performed by killing molecules, such as granzymes and perforin, and Fas-Fas ligand (Fairclough et al., 2008). After recognition of target cells by effector cells, granules released from killer cells move to the immunological synapse by exocytosis. Their membrane then fuses with the killer cell plasma membrane and releases contents into target cells. Previous studies have shown that increased expression of perforin and granzyme B were detected on CD8⁺ T cells from BAL fluid in COPD patients (Hodge et al., 2006). The percentage of both granzyme B and perforin positive NKT-like

and NK cells were significantly higher in induced sputum from COPD patients (Urbanowicz et al., 2010), but lower in peripheral blood samples (Urbanowicz et al., 2009). Morissette and colleague found that the percentage of peripheral CD8⁺ T cells and CD56⁺ cells expressing perforin, granzyme B and Fas-ligand was not different between healthy non-smokers, healthy smokers and emphysematous smokers (Morissette et al., 2007). However, it is hard to conclude that perforin and granzyme B positive NKT cells are the critical difference between COPD and emphysematous conditions. More studies should be carried out to confirm it. Furthermore, up-regulated soluble Fas was observed in plasma from severe COPD patients but not mild or moderate COPD patients (Yasuda et al., 1998).

In PBMCs, IL-15 not only stimulates NK cells or NKT cells, but also other immune cells, such as memory CD4⁺ T cells and CD8⁺ T cells, which consequently secrete more functional cytokines and chemokines to induce adaptive immune response (Anderson et al., 1995, Cosman et al., 1995). To confirm whether the suppression effects caused by CSE were directly related to IL-15 treated NKT-like cells and NK cells, in this section, we isolated NKT-like cells (CD3⁺CD56⁺) and NK cells (CD3⁻CD56⁺) from PBMCs in healthy non-smoker, current

healthy smokers and current smokers with COPD. Both activation levels and killing molecule expression were investigated and compared in three groups. Two types of cells were incubated with medium alone, CSE alone or IL-15 in the absence or presence of CSE, for 48 hours. Cell activation markers CD69 and CD25 were measured, along with killing molecules including perforin, granzyme B and Fas-ligand.

3.3.2 Material and methods

3.3.2.1 Study population and procedures

The criteria of participant enrolments are the same as previously introduced. Nine participants were included in this part of the study. Table 3.3.1 shows the demographic and spirometric data of the participants.

3.3.2.2 Reagents

Recombined human IL-15 was obtained from PeproTech EC (London, UK). Other monoclonal antibodies for flowcytometry were shown in table 3.3.2.

3.3.2.3 Cigarette smoke extract preparation

CSE were freshly prepared prior to 30 minutes of cell culture for each experiment to avoid breakdown of substances in the

Table 3.3.1 Demographic and spirometric values of the CSE suppression study groups

	Healthy non-smokers	Current healthy smokers	cuS-COPD
Participants	3	3	3
Age (years)	52 (50-54)	57 (54-60)	55 (52-58)
Gender (M/F)	0/3	2/1	2/1
Packs/yrs	0	39 (31-42)	43 (35-47)
Smoking status (Current/Ex)	0	3/0	3/0
Chronic bronchitis (Yes/No)	0	2/1	3/0
FEV₁ (% pred)	111 (102-119)	105 (98-117)	52 (46-60)
FEV₁/FVC (%)	97 (86-113)	90 (69-106)	49(45-53)
ΔFEV₁ post bronch	2.2 (1.7-2.5)	2.0 (1.7-2.4)	1.5 (1.2-1.6)
BMI (kg/m²)	25.8 (22.6-28.3)	26.7 (20.5-29.4)	20.3 (15.8-25.1)
Inhaled GCS (on/off)	N/A	N/A	0/3
MRC dyspnoea scale	N/A	N/A	3 (2-4)
Distance walked in 6 min (m)	N/A	N/A	277 (196-312)
BODE Index	N/A	N/A	4 (3-6)

extract and evaporation of volatile components. More details are provided in section 2.4.

3.3.2.4 Separation of NKT-like cells and NK cells

Isolated PBMCs were incubated with anti-CD56 MicroBeads (Miltenyi Biotech Ltd, Surrey, UK) in MACS buffer containing sterile D-PBS (Sigma), 1% heat-inactivated fetal calf serum (FCS) (QB-Perbio) and 0.4% EDTA (Sigma) for 15 minutes at 4°C. More details are provided in section 2.3.

Table 3.3.2 Antibodies used for flow cytometry

Antigen	Fluorochrome	Isotype	Clone	Source
CD3	PC7	Mouse IgG1	UCHT1	Beckman Coulter, Luton, UK
CD25	PE	Mouse IgG2a,	B1.49.9	Beckman Coulter, Luton, UK
CD56	PC5	Mouse IgG1	N901	Beckman Coulter, Luton, UK
CD69	ECD	Mouse IgG2b	TP1.55.3	Beckman Coulter, Luton, UK
KIR3DL1 (CD158e1)	FITC	Mouse IgG1, k	DX9	Biolegend, San Diego, CA, USA
Fas-L (CD178)	PE	IgG Hamster	4A5	Beckman Coulter, Luton, UK
GranzymeB	FITC	Mouse IgG1, k	GB11	BD Pharmingen™
Perforin	PE	Mouse IgG2b, k	δG9	BD Pharmingen™

3.3.2.5 Cell culture and treatments

Isolated NKT-like cells and NK cells were separately resuspended in RPMI-1640 medium supplemented with 10% of FCS at a density of 1×10^6 cells/ml and cultured at 200ul/well in a 96-well flat bottom tissue culture plate. These cells were either untreated or treated with CSE, IL-15 or IL-15 plus CSE. All cells were cultured at 37°C in a 5% CO₂ for 48 hours before analysis.

3.3.2.6 Extracellular staining for CD56⁺ cells

Fluorescently conjugated antibodies used for extracellular staining were as follows, CD3 (PC7), CD56 (PC5), CD69 (ECD) CD25 (PE) and Fas-L (PE), or KIR (FITC). More details are shown in section 2.6.

3.3.2.7 Intracellular staining for CD56⁺ cells

Fluorescently conjugated antibodies including perforin (PE) and granzyme B (FITC) were used for intracellular staining. Briefly, Brefeldin A (Sigma, 0.4ul for per 2×10^5 cells) diluted in RPMI medium was added to cell culture prior to 14 hours before harvesting cells. Cell suspension were washed with PBA, fixed with 3% of formaldehyde fix and incubated in the dark at 4°C for 5 minutes. More details are shown in section 2.6.

3.3.2.8 Flow cytometric analysis

Flow cytometric analysis of antibody labelled cells was performed using an EPICS Altra (Beckman Coulter, Luton, UK). More details are provided in section 2.9.

3.3.2.9 Statistical analysis

The statistical analysis was performed with Prism software, version 4.0c (GraphPad). Differences between the three groups of participants were tested using the parametric

student's *t* test. Confidence interval 99% was used to show two-tailed *p* values. *P* values of less than 0.01 were considered to indicate statistical significance.

3.3.3 Results

3.3.3.1 Demographic and medical characteristics of COPD participants, healthy smokers and healthy controls

There were no statistical differences between groups in terms of age or pack years smoked between current smokers with COPD and current healthy smokers (Data not shown).

3.3.3.2 Effects of CSE on CD56⁺ cell activation

Isolated NKT-like cells and NK cells were incubated with medium alone, CSE (1%) alone, IL-15 (20ug/ml) in the absence or in the presence of CSE (1%) for 48 hours at 37°C before cell collection. Cells are designated activated when they express one or both activation markers (i.e., CD69 positive, CD69/CD25 double positive or CD25 positive). There were no significant differences on the activation levels of NKT-like cells (Figure 3.3.1.A) or NK cells (Figure 3.3.1.C) between medium and CSE alone incubation in all groups (all *p*>0.01). As shown in figure 3.3.1.B, activation levels of CSE treated NKT-like cells upon IL-15 stimulation were significantly suppressed

compared to IL-15 itself mediated cell activation in all groups ($p < 0.01$). Also, CSE very markedly suppressed IL-15 mediated NK cell activation in all groups (Figure 3.3.1.D; HNS, $p < 0.005$; HS, $p < 0.005$; cuS-COPD, $p < 0.001$). Figure 3.3.2 shows the representative dot plots of CSE treated NKT-like cell and NK cell activation upon IL-15 stimulation in current smokers with COPD.

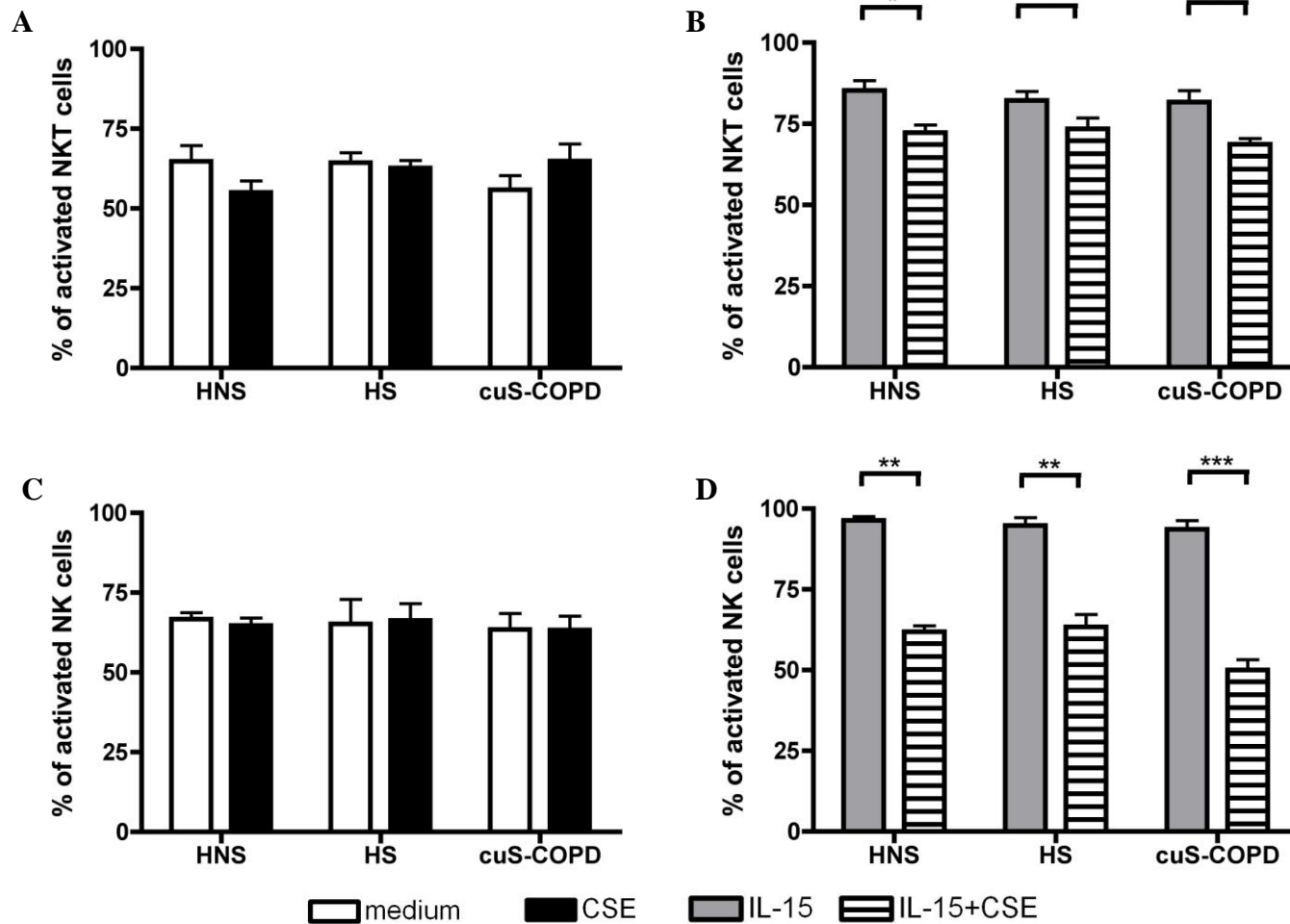


Figure 3.3.1 CSE effects on activation levels of NKT-like cells and NK cells in different conditioned medium. NKT-like cells (panel A and B) and NK cells (panel C and D) were isolated from PBMCs in three groups including health non-smokers (HNS, n=3), current healthy smokers (HS, n=3), and current smokers with COPD (cuS-COPD, n=3). Cells were untreated or treated by CSE alone, IL-15 alone or IL-15+CSE and incubated 48 hours. Significant differences were considered as p value<0.05. *, p<0.01; **, p<0.005; ***, p<0.001. Cells are designated activated when they express one or both activation markers including CD69 and CD25.

3.3.3.3 Effects of CSE on killing molecule expression by CD56⁺ cells

Killing molecules, such as granzymes and perforin, and Fas-Fas ligand, perform major functions in cytotoxic cell mediated immune responses. Here we detected, *in vitro*, the effects of CSE on expression of granzyme B (Figure 3.3.3), perforin (Figure 3.3.4), and Fas-ligand (Figure 3.3.5) in NKT-like cells and NK cells upon IL-15 stimulation. There were no significant differences on the expression of granzyme B by either NKT cells or NK cells that were treated with or without CSE (Figure 3.3.3.A and Figure 3.3.3.C). The expression of granzyme B on IL-15-activated NKT-like cells were significantly inhibited by CSE in healthy non-smokers ($p < 0.01$) (Figure 3.3.3.B). Also, CSE dramatically reduced the expression of granzyme B on IL-15-treated NK cells in both healthy non-smokers (Figure 3.3.3.D, $p < 0.01$) and current smokers with COPD (Figure 3.3.3.D, $p < 0.001$). Perforin was highly expressed in both NKT-like cells and NK cells with or without extra stimulation in all groups (Figure 3.3.4). Whereas Fas-ligand expressed by NKT-like cells and NK cells was lower in all cell incubation conditions in three studied groups (Figure 3.3.5). Table 3.3.3 and table 3.3.4 showed the actual value range of expression levels of molecules of interest on NKT cells and NK cells.

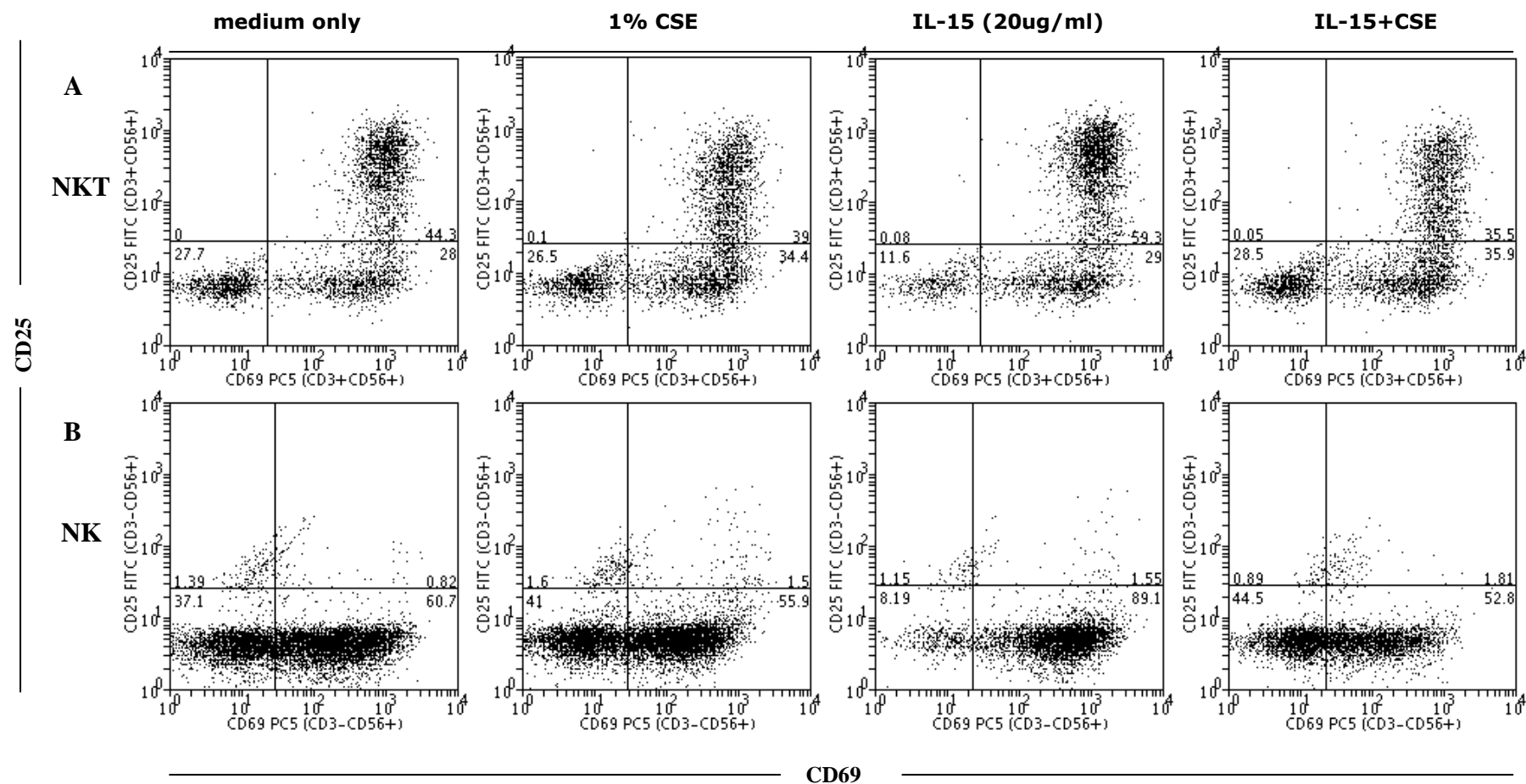
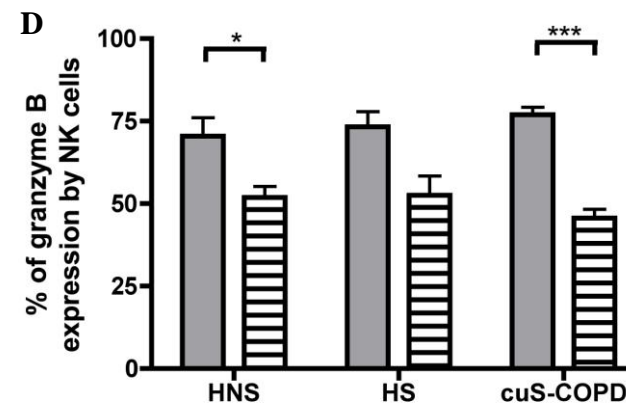
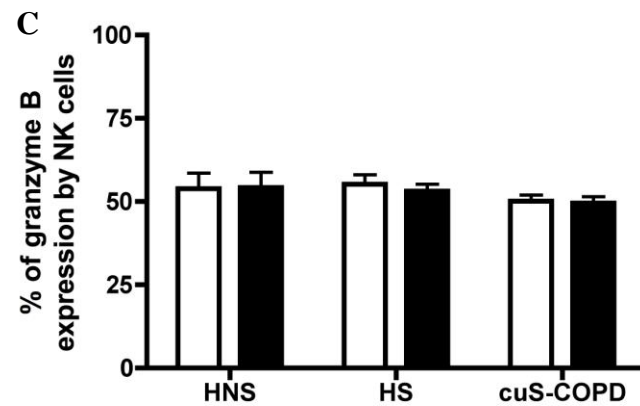
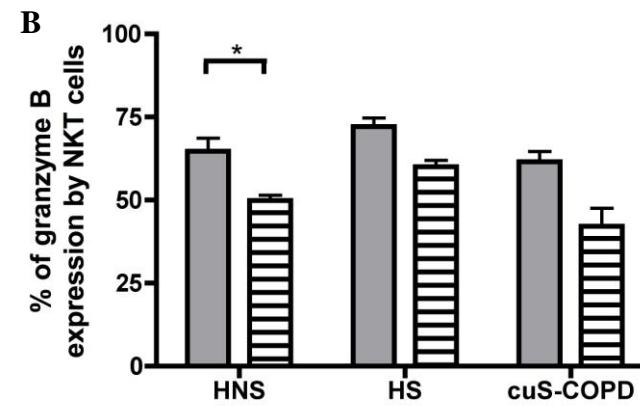
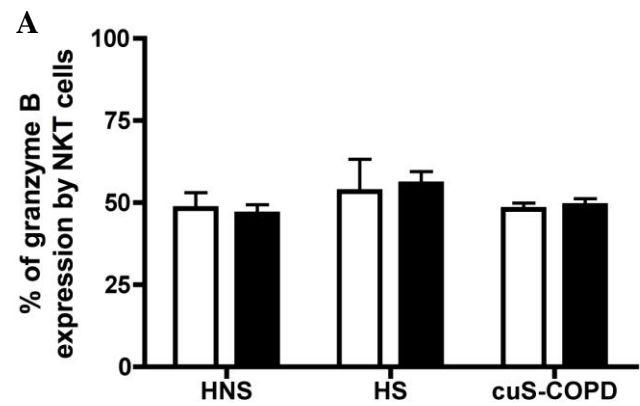


Figure 3.3.2 Representative dot plots for activation of NKT-like cells (panel A) and NK cells (panel B) isolated from PBMCs from a current smoker with COPD. NKT-like cells or NK cells were either untreated or treated with CSE alone, IL-15 alone or IL-15 co-treated with CSE.



medium
 CSE
 IL-15
 IL-15+CSE

Figure 3.3.3 CSE effects on expression of granzyme B by NKT-like cells and NK cells. NKT-like cells (panel A and B) and NK cells (panel C and D) were isolated from PBMCs in three groups including health non-smokers (HNS, n=3), current healthy smokers (HS, n=3), and current smokers with COPD (cuS-COPD, n=3). Cells were untreated or treated by CSE alone, IL-15 alone or IL-15+CSE and incubated 48 hours. *, p<0.01; **, p<0.005; ***, p<0.001.

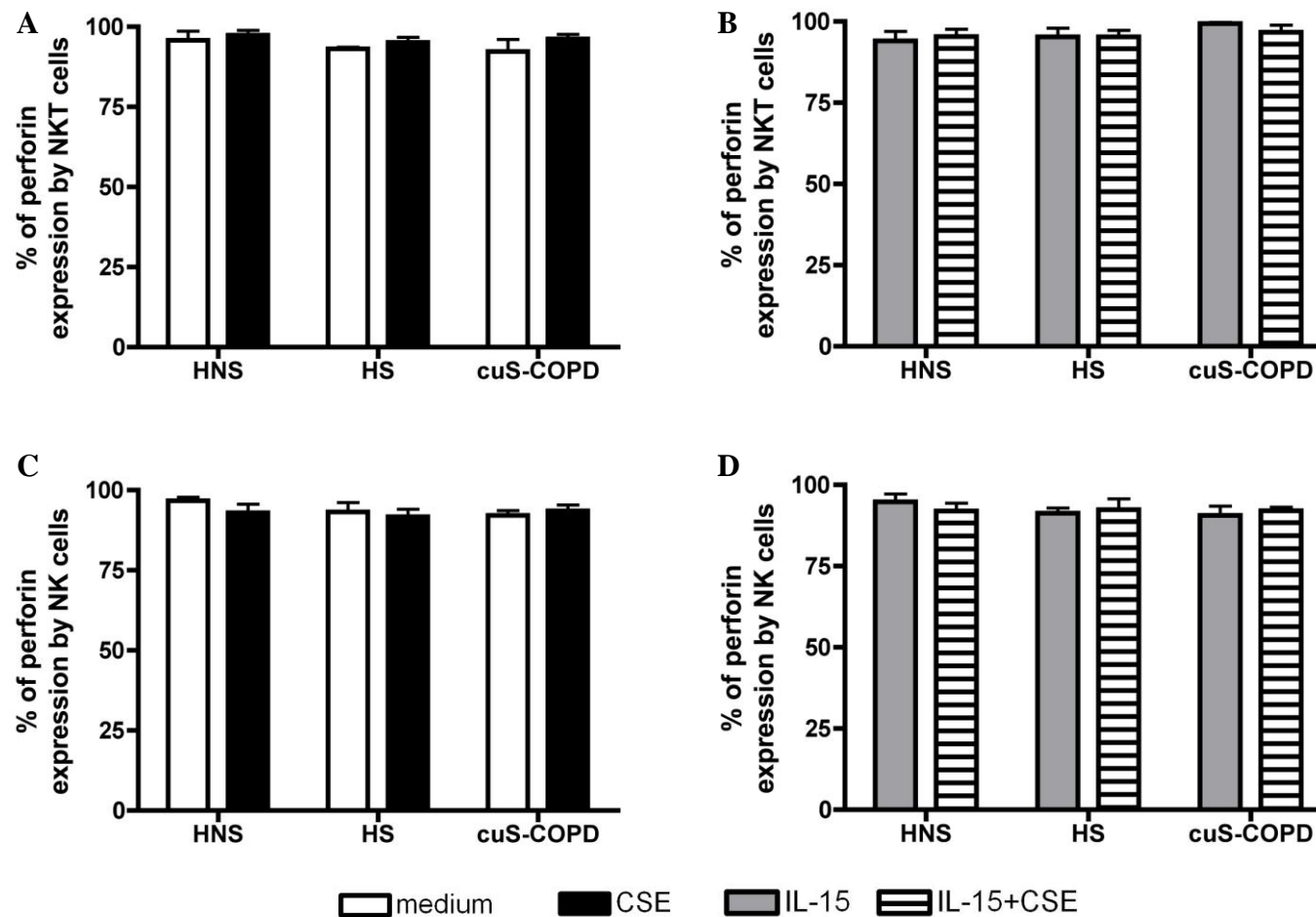


Figure 3.3.4 CSE effects on expression of perforin by NKT-like cells and NK cells. NKT-like cells (panel A and B) and NK cells (panel C and D) were isolated from PBMCs in three groups including health non-smokers (HNS, n=3), current healthy smokers (HS, n=3), and current smokers with COPD (cuS-COPD, n=3). Cells were untreated or treated by CSE alone, IL-15 alone or IL-15+CSE and incubated 48 hours. *, $p < 0.01$; **, $p < 0.005$; ***, $p < 0.001$.

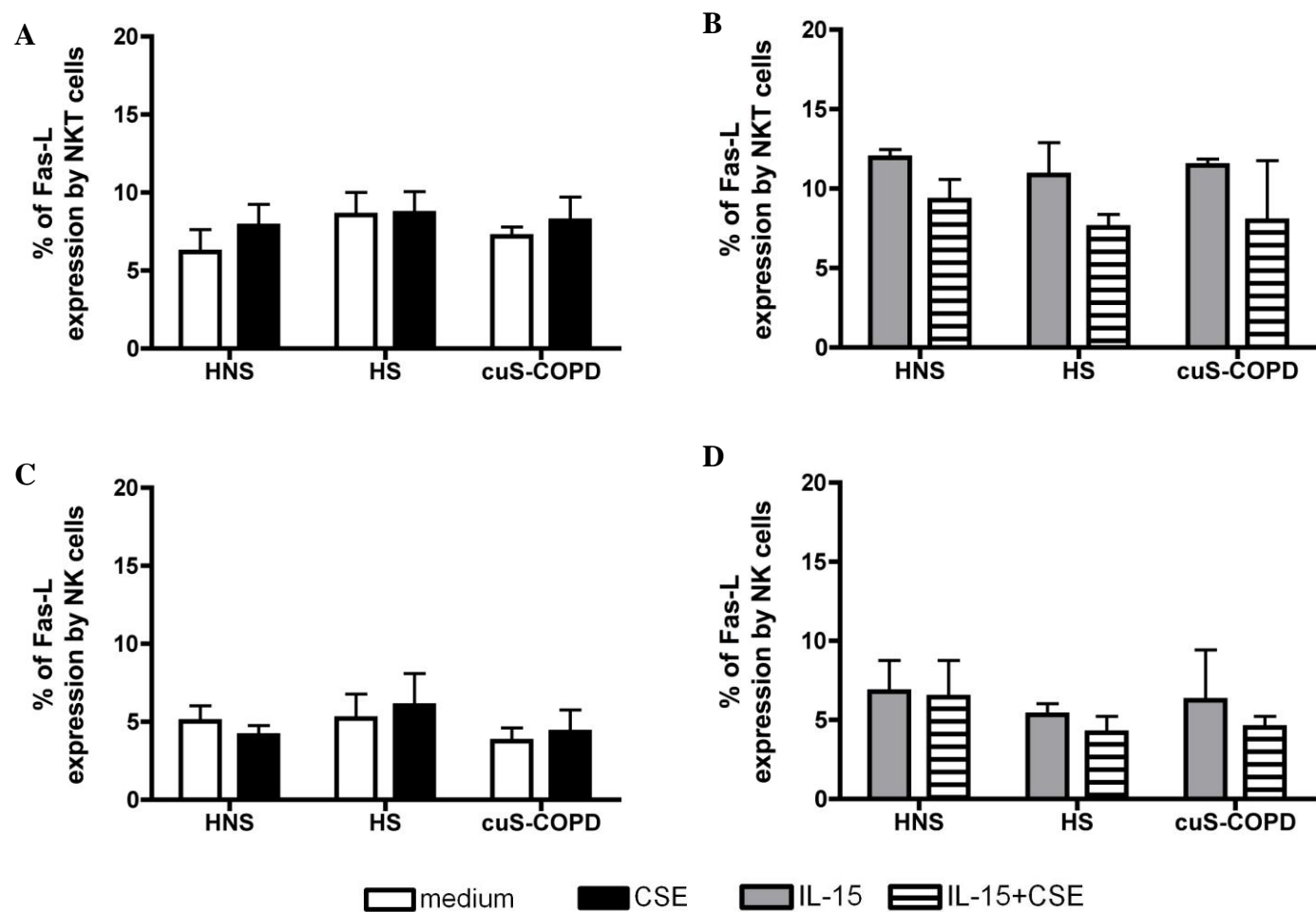


Figure 3.3.5 CSE effects on expression of Fas-ligand by NKT-like cells and NK cells. NKT-like cells (panel A and B) and NK cells (panel C and D) were isolated from PBMCs in three groups including health non-smokers (HNS, n=3), current healthy smokers (HS, n=3), and current smokers with COPD (cuS-COPD, n=3). Cells were untreated or treated by CSE alone, IL-15 alone or IL-15+CSE and incubated 48 hours. *, $p < 0.01$; **, $p < 0.005$; ***, $p < 0.001$.

Table 3.3.3 Expression of molecules of interest on NKT cells with or without activation.

Granzyme B				
	Medium only	CSE only	IL-15	IL-15 + CSE
HNS	48.2% (39.2-55.6)	46.6% (43.6-52.1)	64.7% (57.8-71.3)	49.8% (47.6-52.9)
HS	53.3% (37.8-71.6)	55.7% (50.0-62.0)	72.1% (69.1-77.2)	60.1% (56.8-63.3)
CuS-COPD	47.9% (44.0-50.0)	49.1% (45.1-52.2)	61.5% (55.7-66.6)	42.1% (33.7-52.2)
Perforin				
	Medium only	CSE only	IL-15	IL-15 + CSE
HNS	95.8% (90.1-99.1)	97.3% (94.2-99.2)	94.0% (90.3-99.8)	95.4% (91.2-98.8)
HS	93.1% (92.3-94.0)	95.1% (92.2-97.6)	95.3% (90.3-99.0)	95.3% (91.7-98.8)
CuS-COPD	92.3% (87.2-99.6)	96.2% (93.3-98)	99.31% (98.2-99.9)	96.7% (92.4-99.0)
Fas-ligand				
	Medium only	CSE only	IL-15	IL-15 + CSE
HNS	6.2% (3.4-8.0)	7.9% (5.6-10.3)	12.0% (11.0-12.7)	9.3% (7.2-11.7)
HS	8.6% (5.7-10.2)	8.7% (6.6-11.3)	10.9% (7.3-14.4)	7.6% (6.3-9.1)
CuS-COPD	7.2% (6.1-8.1)	8.2% (6.4-11.2)	11.5% (11.0-12.3)	8.0% (2.1-15.1)
KIR				
	Medium only	CSE only	IL-15	IL-15 + CSE
HNS	7.9% (6.0-9.9)	8.3% (5.1-10.1)	9.0% (5.1-11.2)	8.5% (7.3-9.3)
HS	9.8% (7.4-12.1)	9.3% (8.8-10.3)	11.2% (8.2-15.2)	7.3% (5.5-9.2)
CuS-COPD	9.0% (8.9-9.1)	9.7% (8.0-11.2)	12.7% (10.3-15.1)	6.8% (2.5-10.2)

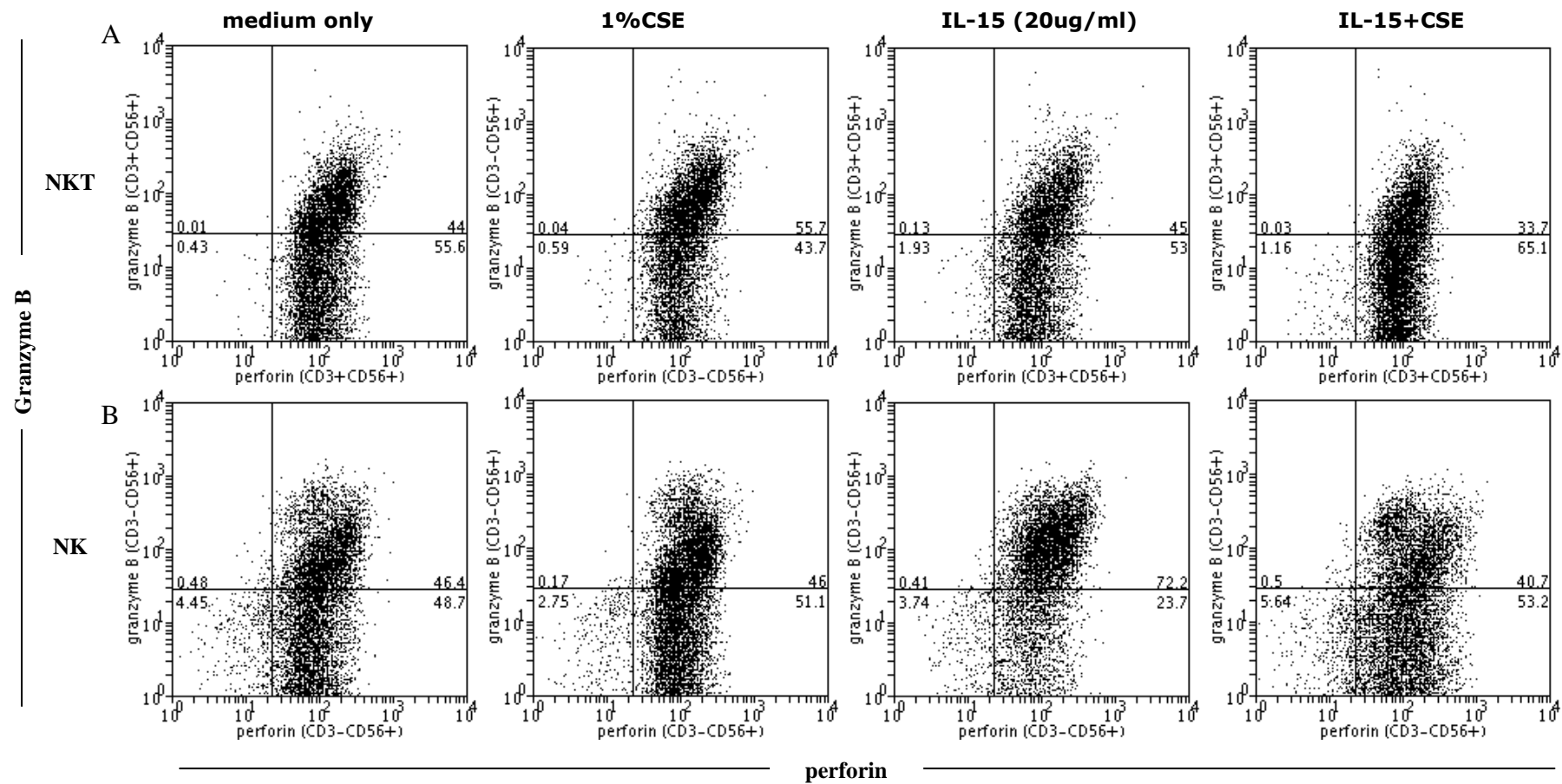
Table 3.3.4. Expression of molecules of interest on NK cells with or without activation.

Granzyme B				
	Medium only	CSE only	IL-15	IL-15 + CSE
HNS	53.9% (44.9-60.1)	54.2% (45.2-59.3)	70.5% (61.1-80.3)	51.9% (46.5-57.9)
HS	55.2% (50.1-60.1)	53.1% (49.0-56.9)	73.3% (67.0-82.2)	52.6% (42.0-62.1)
CuS-COPD	50.1% (46.9-53.4)	49.6% (46.2-52.6)	76.9% (72.6-80.3)	45.7% (41.2-50.2)
Perforin				
	Medium only	CSE only	IL-15	IL-15 + CSE
HNS	96.6% (95.5-98.7)	92.9% (90.0-98.3)	94.8% (90.0-97.7)	91.9% (87.3-95.5)
HS	93.2% (91.0-99.2)	91.7% (87.9-95.9)	91.3% (89.4-94.4)	92.4% (88-98.9)
CuS-COPD	92.1% (90.2-95.1)	93.6% (90.9-97.1)	90.6% (85.6-95.5)	92.1% (90.1-93.9)
Fas-ligand				
	Medium only	CSE only	IL-15	IL-15 + CSE
HNS	5.0% (3.8-7.0)	4.1% (3.0-5.1)	12.0% (3.4-10.2)	9.3% (3.4-11.0)
HS	5.2% (2.3-7.7)	6.0% (2.9-9.9)	10.9% (3.9-6.2)	7.6% (2.4-5.9)
CuS-COPD	3.8% (2.2-5.1)	4.3% (2.1-7)	11.5% (0.3-11.2)	8.0% (3.2-5.6)
KIR				
	Medium only	CSE only	IL-15	IL-15 + CSE
HNS	5.8% (3.3-7.9)	4.3% (0.9-9.1)	5.0% (0.7-8.9)	5.4% (2.5-10.2)
HS	4.6% (2.1-7.2)	3.1% (1.8-5.4)	5.3% (2.9-7.6)	4.3% (2.4-6.0)
CuS-COPD	3.4% (1.5-5.7)	2.1% (0.7-3.5)	6.1% (0.6-10.1)	2.6% (1.7-3.3)

CSE had no effects on the expression levels of either perforin or Fas-ligand. Collectively, IL-15 had the ability of inducing more granzyme B expression on CD56⁺ cells but had no effects on the expression of perforin and Fas-ligand. Figure 3.3.6 shows representative dot plots of CSE effects on killing molecule expression by CD56⁺ cells from current smokers with COPD.

3.3.3.4 Effects of CSE on KIR expression by CD56⁺ cells

The effect of CSE on KIR expression was also studied (Figure 3.3.7). Results in chapter 3.1 demonstrated that the expression of KIR (CD158e1), *in vivo*, was much more smoking-related than disease related (Figure 3.1.8). However, here, *in vitro*, we did not detect any statistical differences on KIR expression by CSE treated CD56⁺ cells upon IL-15 stimulation KIR expression showed very low levels on CD56⁺ cells in each group. Due to low cell numbers of CD56⁺ cells, we did not detect the NKG2D expression.



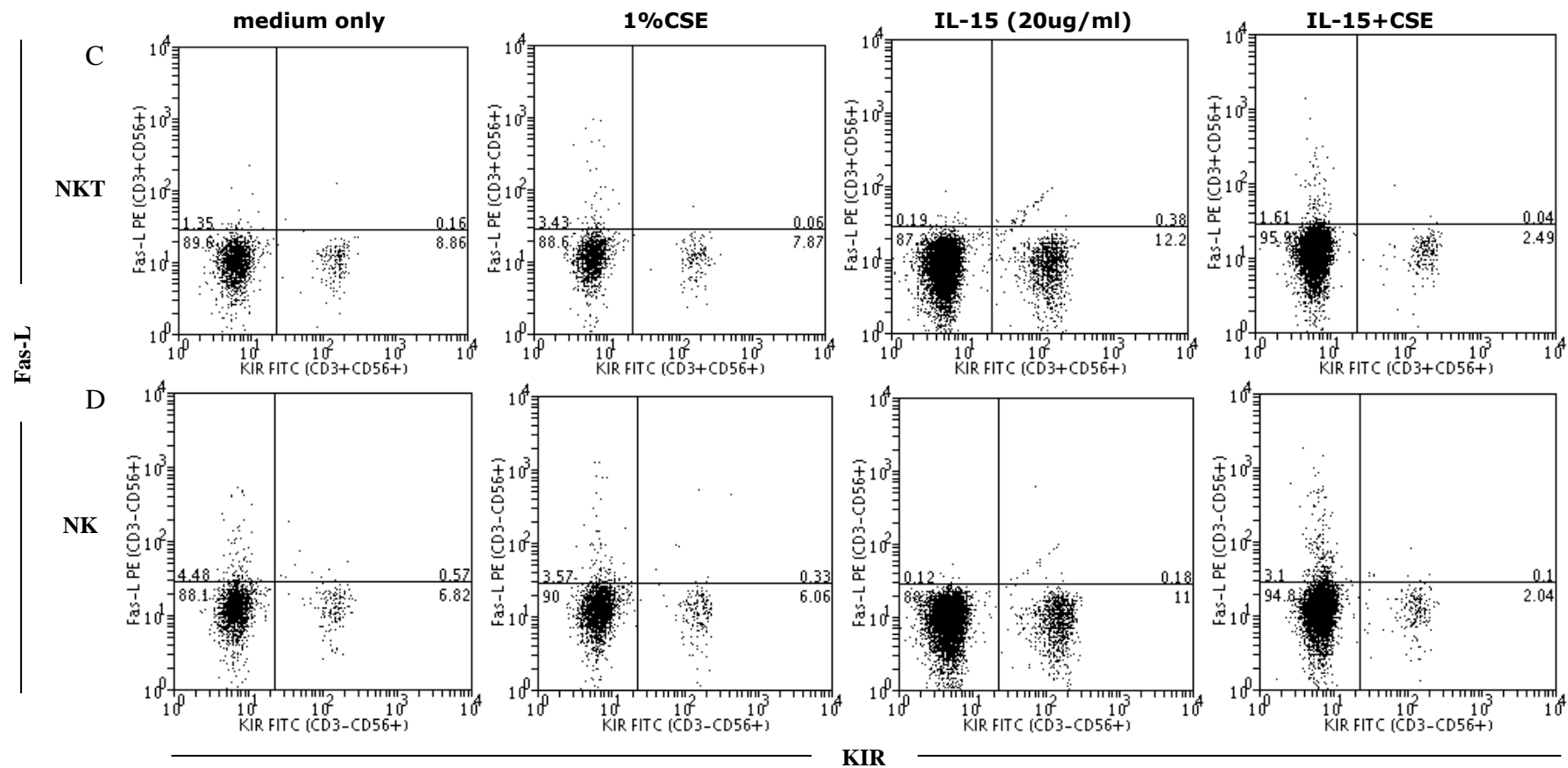


Figure 3.3.6 Representative dot plots for expression of perforin, granzyme B, Fas-ligand and KIR by NKT-like cells (panel A and B) and NK cells (panel C and D) isolated from PBMCs from a current smoker with COPD. NKT-like cells or NK cells were either untreated or treated with CSE alone, IL-15 alone or IL-15 co-treated with CSE.

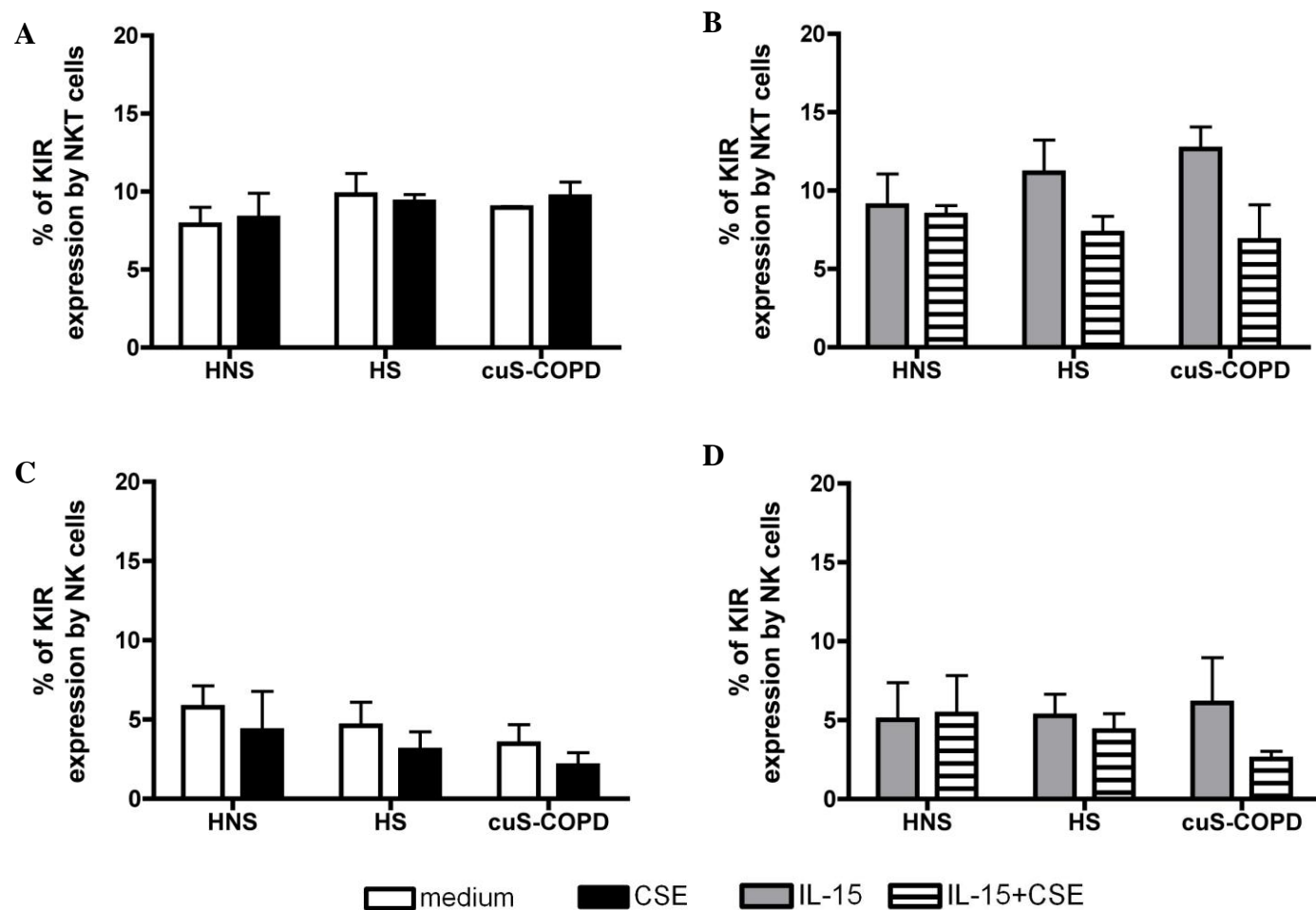


Figure 3.3.7 CSE effects on expression of KIR by NKT-like cells and NK cells. NKT-like cells (panel A and B) and NK cells (panel C and D) were isolated from PBMCs in three groups including health non-smokers (HNS, n=3), current healthy smokers (HS, n=3), and current smokers with COPD (cuS-COPD, n=3). Cells were untreated or treated by CSE alone, IL-15 alone or IL-15+CSE and incubated 48 hours. *, $p < 0.01$; **, $p < 0.005$; ***, $p < 0.001$.

3.3.3.5 CSE suppressed cell activation

As previously mentioned, cell activation was measured by expressing one or both of the activation markers including CD69 and CD25. CSE significantly suppressed the activation levels of IL-15 treated NKT-like cells and NK cells. Here we compared the absolute suppression induced by CSE on IL-15 mediated NKT-like cell or NK cell activation in three groups. There were no statistical differences on the suppression levels between groups (Figure 3.3.8.A and B). The expression levels of granzyme B on CSE treated NKT-like cells or NK cells in response to IL-15 were also compared. No significant differences were detected on the absolute suppression of granzyme B expression between groups (Figure 3.3.9.A and B).

3.3.4 Discussion

Here we report, for the first time, *in vitro*, that CSE significantly suppresses the activation levels of IL-15 treated NKT-like cells and NK cells isolated from PBMCs in healthy non-smokers, current healthy smokers and current smokers with COPD.

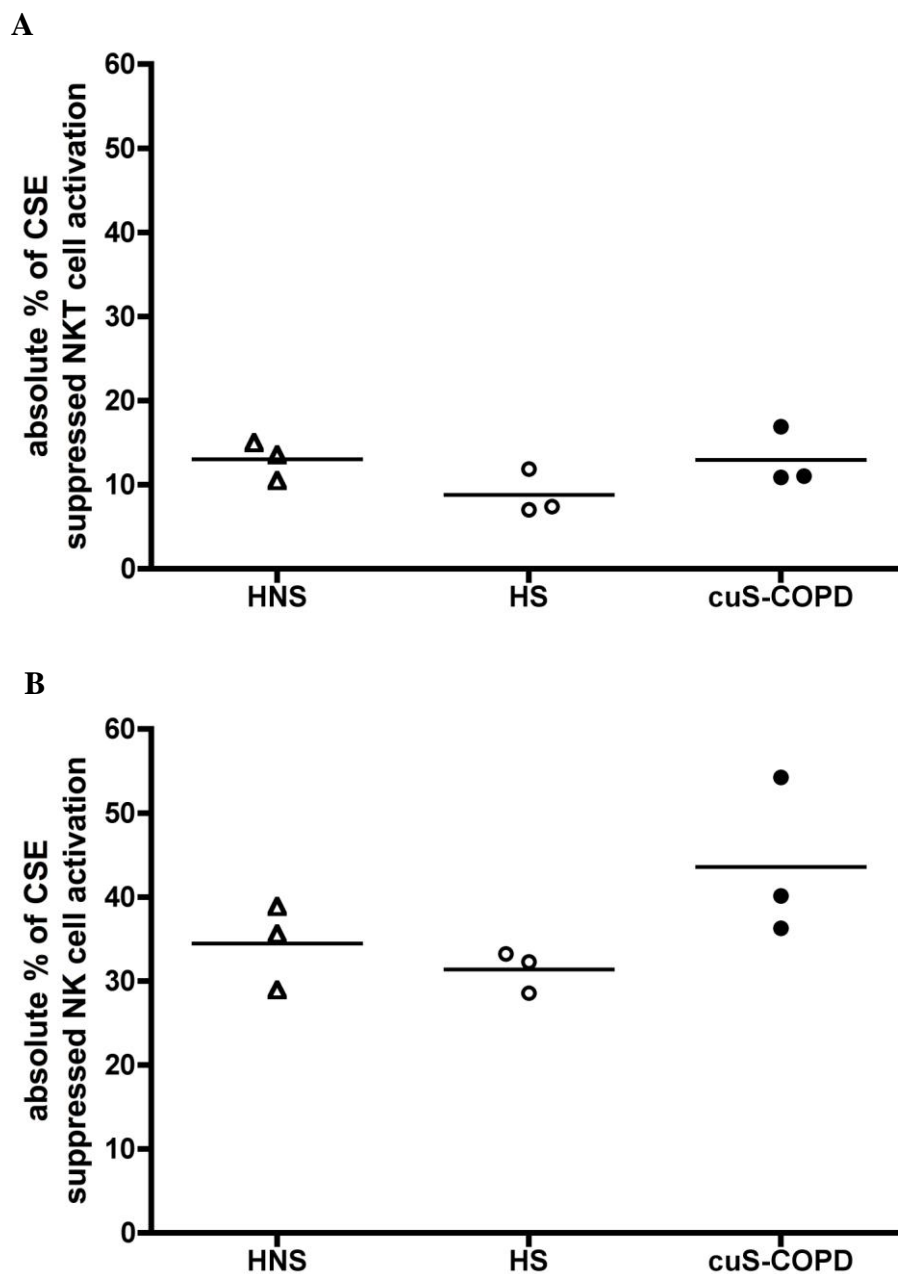


Figure 3.3.8 CSE induced suppression on activation levels of IL-15 stimulated NKT-like cells and NK cells in three groups. Absolute suppression levels were gained by using (% of IL-15 activated cells) minus (% of IL-15+CSE treated cells). All p value>0.05. HNS (healthy non-smokers, n=3); HS (healthy smokers, n=3); cuS-COPD (current smokers with COPD, n=3).

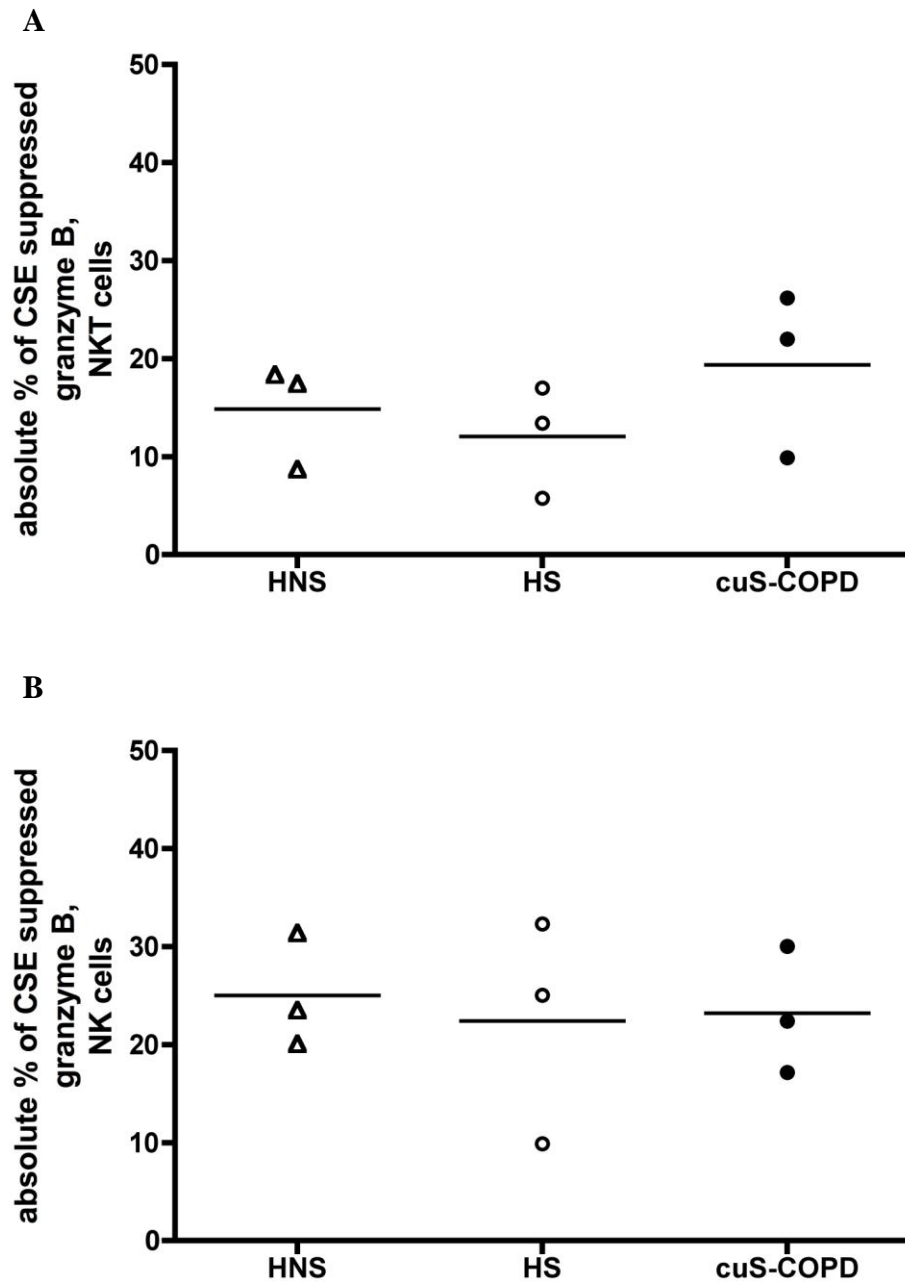


Figure 3.3.9 CSE induced suppression on expression of granzyme B by IL-15 stimulated NKT-like cells and NK cells in three groups. Absolute suppression levels were gained by using (% of IL-15 activated cells) minus (% of IL-15+CSE treated cells). All $p > 0.05$. HNS (healthy non-smokers, $n=3$); HS (healthy smokers, $n=3$); cuS-COPD (current smokers with COPD, $n=3$).

IL-15 is a potent cytokine with the ability of stimulating many kinds of immune cells, such as memory CD4⁺ T cells, naive or memory CD8⁺ T cells, DCs, NKT cells and NK cells, which can further secrete more cytokines and chemokines to induce adaptive immune responses (Anderson et al., 1995, Cosman et al., 1995). Several studies demonstrated that cigarette smoke inhibited IL-12 and type-1 IFN production in activated PBMCs, which consequently decreased the IL-15 levels in healthy non-smokers (Mian et al., 2008, Mian et al., 2009). Also, both *in vitro* and *in vivo*, cigarette smoke significantly diminished IL-15 mRNA levels in poly I:C-activated PBMCs from healthy non-smoking participants. IL-15 production was markedly impaired in smoke-conditioned medium treated PBMCs upon poly I:C simulation. In result chapter 3.2, we showed that the CSE significantly reduced the activation levels of NKT-like cells from IL-15 treated PBMCs in healthy volunteers and ex-smokers with COPD. Also, CSE dramatically decreased the activation levels of NK cells in all groups, especially in COPD patients. Therefore, we examined whether the suppression induced by CSE on CD56⁺ cells is directly correlated with IL-15 or through other immune mediators secreted by other cells. We found that CSE itself did not show any statistical impacts on the activation levels of CD56⁺ cells compared to untreated cells. IL-15 highly activated NKT-like

cells and NK cells in both healthy volunteers and COPD patients. However, when CSE was added in the cultures, their activation levels were significantly reduced. This effect on NK cells was much more significant in current smokers with COPD. Interestingly, after 48 hour incubation, activation stages of NKT cells and NK cells showed different status (Figure 3.3.2). Most activated NKT cells are both CD69 and CD25 positive, whereas NK cells are mainly CD69 positive. This might be explained as followed. Cytokines released by NKT cells are always rapid and transient, with a peak of activity within 24 hours. After 24 hours, productions of these cytokines, for example IFN- γ , quickly return to baseline levels (Eberl and MacDonald, 1998). IFN- γ is a good inducer for NK cell activation. This may explain why NKT cells were more quickly activated than NK cells.

Our group has illustrated that the cytotoxic functions of NK cells and NKT cells from PBMCs were both markedly reduced in COPD patients compared to healthy volunteers (Urbanowicz et al., 2009). *In vitro*, the percentage of perforin /granzyme B double positive NK cells was significantly lower in COPD patients compared to controls. Furthermore, Mian *et al* demonstrated that cigarette smoke inhibited IL-15 mediated NK cell cytotoxic killing ability with expressing lower NK cell

degranulation marker CD107a in healthy non-smokers (Mian et al., 2009). Granzymes, perforin and Fas-Fas ligand are three major killing molecules that facilitate NKT-like cells and NK cells performing their cytotoxic functions. To evaluate whether CSE induced suppression was linked to NK cell cytotoxicity, we analysed the expression levels of above three cytotoxic granules on IL-15 stimulated CD56⁺ cells in the presence or in the absence of CSE. Surprisingly, CSE did not induce any remarkable impacts on expression levels of perforin and Fas-ligand in all groups. Perforin was highly expressed in all stimulating conditions whereas Fas-L showed very low levels. Of note, expression of granzyme B on IL-15 treated NKT-like cells was markedly reduced by CSE in healthy non-smokers. The remarkable effects of CSE on granzyme B expression by NK cells were also detectable in healthy non-smokers and current smokers with COPD. These findings were partially supported by other research. Morissette and colleague illustrated that no statistical differences were observed on the expression of perforin and granzyme B in peripheral NK cells from either healthy volunteers or smokers with evidence of emphysema. Protein expression of Fas-ligand was not detected by flow cytometry (Morissette et al., 2007). Although there was a markedly increase in soluble Fas in plasma from severe COPD patients, there were no differences

on its expression in mild or moderate COPD and healthy controls (Yasuda et al., 1998). Hodge *et al* also demonstrated that there were no significant changes in the expression levels of granzyme A by NK or NKT-like cells in the blood of COPD patients compared to controls (Hodge et al., 2013). Taken together, our data suggest that granzyme B, but not perforin and Fas-ligand, was a functional molecule attributing to the significant impairment NK cell killing ability in response to CSE in COPD patients.

KIRs are inhibitory receptors expressed by cytotoxic cells. We early found that, *in vivo*, expression levels of KIR (CD158e1) by killer cells including CD8⁺ T cells and CD56⁺ cells were negatively correlated with the activation levels of these cells in COPD patients. In this study, KIR expression by IL-15 incubated CD56⁺ cells in response to CSE was analysed and compared between groups. Here, *in vitro*, we first time showed that there were no significant differences on KIR (CD158e1) expression by CD56⁺ in different cell culture conditions. KIR expression was at very low levels in each group. Few studies were carried out to detect the KIRs expression in either healthy volunteers or COPD patients. However, one study reported that NKG2D expression was not inhibited in cigarette smoke treated PBMCs after poly I:C

induction (Mian et al., 2009). NKG2D is an activating receptor mainly expressed by killer cells and γ/δ T cells. We found, *ex vivo*, it was highly expressed in CD56⁺ cells (averagely over 90%) in both healthy participants and COPD patients. Collectively, KIR (CD158e1) was not a functional inhibitory receptor for CSE induced suppression of CD56⁺ cells activation in response to IL-15. Also, KIR levels from untreated cells *in vitro* (Figure 3.3.7) were lower than that *ex-vivo* (Fig 3.1.8). These might be due to, *in vitro*, cell interaction during 48 hour incubation. Activated cells might produce inflammatory molecules that could affect the functional receptor expression levels. However, more research should be carried out to clarify it.

Although CSE significantly suppressed CD56⁺ cell activation and their granzyme B expression in IL-15 conditioned medium, there were no statistical differences between groups. This might be due to small sample size used in this study. We indeed found that the activation levels of NK cells inhibited by CSE in COPD patients were much more significant ($p < 0.001$).

Several limitations should be considered in our study. Firstly, only small number of subjects in each group was recruited. Secondly, patient samples were only from current smokers

with COPD but not ex-smokers. Moreover, there are many cytotoxic granules such as granzyme A and granzyme K related to killing mechanisms of CD56⁺ cells. However, proportions of NKT-like cells and NK cells are very low in PBMCs. We could just detect representative killing molecules (granzyme B, perforin and Fas-ligand) and one type of inhibitory receptors (KIR3DL, CD158e1) in our research due to cell numbers. The activation of CD56⁺ cells is regulated by signal balance between inhibitory receptors and activating receptors. We could not present any significant differences about KIR expression on cells between groups (Figure 3.3.7). In future, other inhibitory or activating receptors, such as NKG2D and CD94, could be detected to find out whether they play potent roles in CSE suppressed activation.

In summary, this study showed for the first time that, CSE markedly suppressed IL-15 induced NKT-like cell activation and NK cell activation in healthy volunteers and current smokers with COPD. The granzyme B expression was also impaired on both cell types. Expression of perforin, Fas-ligand and KIR did not vary on CD56⁺ cells either in response to IL-15 or co-stimulated CSE, in all groups. Our findings support the concept that cigarette smoke extract affects the cytotoxic functions of NKT-like and NK cell in COPD patients.

3.4 Cytotoxicity of CD8⁺CD56⁻ T lymphocytes in COPD

3.4.1 Introduction

Cytotoxic T lymphocytes (Tc cells, CD3⁺CD8⁺CD56⁻) express the $\alpha\beta$ T cell receptor. The predominant function of Tc cells is to kill infected cells via producing cytokines, such as IFN- γ , and killing molecules like perforin and granzymes (Henkart, 1985, Zagury, 1982). Binding of Fas-ligand presented by Tc cells to Fas expressed by target cells can also induce apoptosis of infected cells (Nagata and Golstein, 1995). Abundant studies have shown abnormalities of Tc cells in COPD, such as in large airways sub-epithelium (O'Shaughnessy et al., 1997, Lams et al., 2000), peripheral airway (Saetta et al., 1998), parenchyma (Saetta et al., 1999), smooth muscle (Baraldo et al., 2004), bronchial arteries (Peinado et al., 1999, Saetta et al., 1999, Turato et al., 2002), peripheral blood (Urbanowicz et al., 2009), sputum samples (Urbanowicz et al., 2010), and small airways such as bronchoalveolar lavage (BAL) (Saetta et al., 2002, Saetta et al., 2003).

Cell-contact-dependent cytotoxicity is the hallmark of killer cell function (Russell and Ley, 2002). Cytotoxic T cells kill their

targets by programming them to undergo apoptosis. The elimination of infected cells without the destruction of healthy tissue requires the cytotoxic mechanisms of CD8⁺ T cells to be both powerful and accurately targeted. Currently, abundant studies have investigated the cytotoxicity of killer cells, such as NK cells and CD8⁺ T lymphocytes, from cancer patients (Jewett and Tseng, 2013, Levy et al., 2013, Maher and Davies, 2004). However, studies related to COPD are rare. Our group early demonstrated that cytotoxicities of NKT-like cells and NK cells were lower in PBMCs but higher in induced sputum from COPD patients compared to healthy volunteers (Urbanowicz et al., 2009, Urbanowicz et al., 2010). Chrysofakis *et al* also observed increased cytotoxic activity of Tc cells in induced sputum of COPD patients (Chrysofakis et al., 2004). In this chapter, we evaluated the cytotoxic activity of blood CD8⁺CD56⁻ T cells against the P815 cell line in four groups including healthy non-smokers, current healthy smokers, current smokers with COPD and ex-smokers with COPD.

3.4.2 Material and methods

3.4.2.1 Study population and procedures

The criteria of participant enrolments are the same as previously introduced. There were thirteen participants

included in this study. Table 3.4.1 shows the demographic and spirometric data of the participants.

3.4.2.2 PBMCs isolation

PBMCs were isolated from whole blood on a discontinuous Histopaque density gradient (Sigma). More details are provided in section 2.2.

3.4.2.3 CD8⁺CD56⁻ T lymphocytes separation

Isolated PBMCs were incubated with anti-CD56 MicroBeads (Miltenyi Biotech Ltd, Surrey, UK) in MACS buffer containing sterile D-PBS (Sigma), 1% heat-inactivated fetal calf serum (FCS) (QB-Perbio) and 0.4% EDTA (Sigma) for 15 minutes at 4°C. CD56⁻ cell fractions were obtained by cell suspension passing through a cold MS column attached to a magnet. Isolated CD56⁻ fractions were washed and counted. To obtain CD8⁺CD56⁻ T lymphocytes, cells were incubated with Multisort Release Reagent for 10 minutes at 4°C, washed and then incubated with MultiSort Stop Reagent and α -CD8 microbeads for 15 minutes at 4°C. Finally the labelled cells were separated on a refrigerated LS column. Detailed protocol for cell separation was shown on manufacturer's instructions. All isolated fractions were washed, counted and purity confirmed by flow cytometry ($\geq 90\%$).

Table 3.4.1 Demographic and spirometric value of the CSE suppression study groups

	Healthy non-smokers	Current healthy smokers	cuS-COPD	exS-COPD
Participants	4	4	3	2
Age (years)	55 (50-60)	54 (51-60)	55 (52-58)	49 (46-52)
Gender (M/F)	1/3	2/2	2/1	0/2
Packs/yrs	0	39 (33-42)	43 (35-47)	44 (39-49)
Smoking status (Current/Ex)	0	4/0	3/0	0/2
Chronic bronchitis (Yes/No)	0	3/1	3/0	2/0
FEV₁ (% pred)	117 (102-128)	103 (92-117)	52 (46-60)	57 (54-60)
FEV₁/FVC (%)	97 (86-116)	87 (66-106)	49(45-53)	47 (39-56)
ΔFEV₁ post bronch	2.4 (1.7-2.9)	1.8 (1.5-2.4)	1.5 (1.2-1.6)	1.4 (1.3-1.6)
BMI (kg/m²)	26.6 (22.6-28.3)	25.2 (19.7-29.4)	20.3 (15.8-25.1)	20.1 (18.7-21.5)
Inhaled GCS (on/off)	N/A	N/A	0/3	1/1
MRC dyspnoea scale	N/A	N/A	3 (2-4)	3 (2-4)
Distance walked in 6 min (m)	N/A	N/A	277 (196-312)	302 (218-386)
BODE Index	N/A	N/A	4 (3-6)	5 (4-7)

exS-COPD, ex-smokers with COPD; cuS-COPD, current smokers with COPD; FEV₁, forced expiratory volume in 1 second; pre, predicted value; FVC, forced vital capacity; BMI, body mass index; GCS, corticosteroid; MRC, medical research council; BODE, **B**MI, **O**airflow **O**bstruction, **D**yspnoea and **E**xercise capacity.

3.4.2.4 Cell culture

For cytotoxicity assay, isolated CD8⁺CD56⁻ T cells were incubated for 24 hours at 37°C in 5% CO₂. Before cell incubation, the plate was coated with 2µg/ml anti-CD3 (in 50ul D-PBS solution each well) for a minimum of 2 hours at 37°C in 5% CO₂. Target cells (P815 cells) were grown in RPMI-1640 medium supplemented with 10% FCS at 37°C in 5% CO₂ and ready for use when they reached confluence.

3.4.2.5 CD8⁺CD56⁻ T lymphocytes Redirected Cytotoxicity Assay

The cytotoxicity of CD8⁺CD56⁻ T lymphocyte was measured with colorimetric quantification of lactate dehydrogenase LDH released from lysed target cells (cell line P815). More details are provided in section 2.7.

3.4.2.6 Flow cytometric analysis

Flow cytometric analysis of antibody labelled cells was performed using an EPICS Altra (Beckman Coulter, Luton, UK). More details are provided in section 2.9.

3.4.2.7 Statistical analysis

The statistical analysis was performed with Prism software, version 4.0c (GraphPad). Data were non-normally distributed

all are expressed as median and range, unless otherwise stated. Differences between the four groups of participants were tested using parametric student's *t* test. Confidence interval 99% was selected to show two-tailed *p* values. *P* values of less than 0.01 were considered to indicate statistical significance.

3.4.3 Results

3.4.3.1 Purity of Tc lymphocytes

The purity of isolated CD8⁺CD56⁻ T lymphocytes was determined by flow cytometry. Figure 3.4.1.A showed the representative dot plots of the proportion of CD3⁺CD8⁺CD56⁻ T cells in each group, namely, healthy non-smokers (HNS, A i), current healthy smokers (HS, A ii), current smokers with COPD (cuS-COPD, A iii) and ex-smokers with COPD (exS-COPD, A iv). The purity of the immunomagnetically selected CD3⁺CD8⁺ T lymphocytes in isolated CD8⁺CD56⁻ fractions was over 90% (Figure 3.4.1.B).

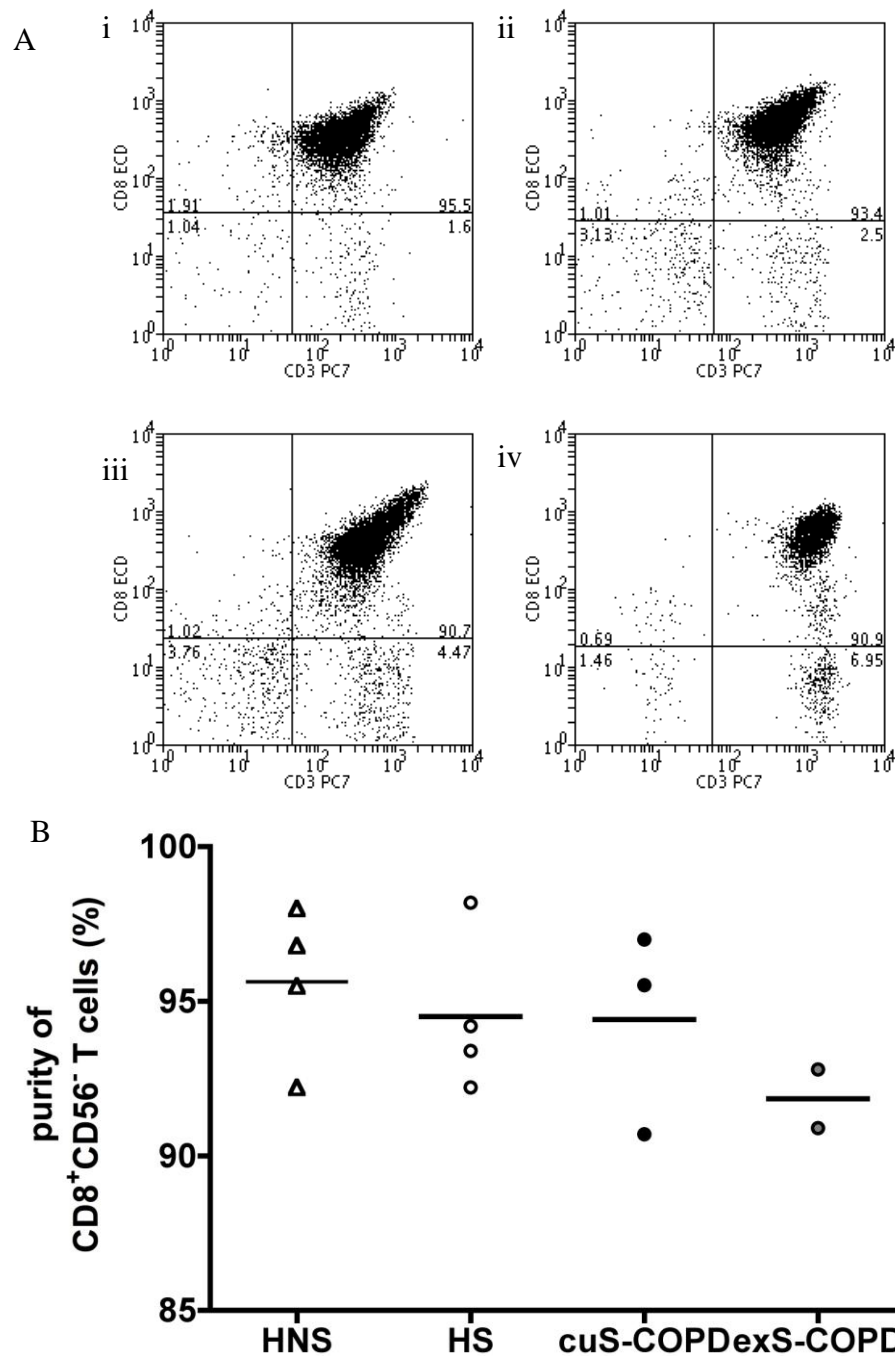


Figure 3.4.1 Purity of immunomagnetically separated $CD8^+CD56^-$ T lymphocytes from the peripheral blood of the studied groups. $CD8^+CD56^-$ T cells were isolated from PBMCs in four groups including health non-smokers (HNS, $n=4$), current healthy smokers (HS, $n=4$), current smokers with COPD (cuS-COPD, $n=3$), and ex-smokers with COPD (exS-COPD, $n=2$). (A) Representative dot plots of the purified $CD3^+CD8^+CD56^-$ T cells from PBMCs in each group, namely, i, NHS; ii, HS; iii, cuS-COPD; iv, exS-COPD. Percentage of purity was shown. (B) Purity of immunomagnetically separated $CD8^+CD56^-$ T lymphocytes in each group.

3.4.3.2 Cytotoxicity of CD8⁺CD56⁻ T lymphocytes in COPD

To test the cytotoxic activity of the immunomagnetically-purified CD8⁺CD56⁻ T cells, a non-radioactive colorimetric quantification of LDH released assay was performed (LDH is a stable cytosolic enzyme released upon cell lysis). *In vitro*, we compared the killing ability of CD8⁺CD56⁻ T cells against the P815 cell line at different ratios in the four participant groups. As shown in Figure 3.4.2, a dose dependent reduction was displayed on the cytotoxic ability of CD8⁺CD56⁻ T cells against P815 cells in each group. At ratio 40:1 (Figure 3.4.3.A), the cytotoxicity of CD8⁺CD56⁻ T cells was significantly less in both current smokers with COPD (21.6% ± 2.4%, $p < 0.005$) and ex-smokers with COPD (27.57% ± 3.6%, $p < 0.005$) compared to healthy non-smokers (63.47% ± 6.6%). The lysis ability of CD8⁺CD56⁻ T cells were also dramatically decreased in current smokers with COPD (21.6% ± 2.4%, $p < 0.005$) and ex-smokers with COPD (27.57% ± 3.6%, $p < 0.01$) compared to current healthy smokers (56.81% ± 9.7%) (Figure 3.4.3.A).

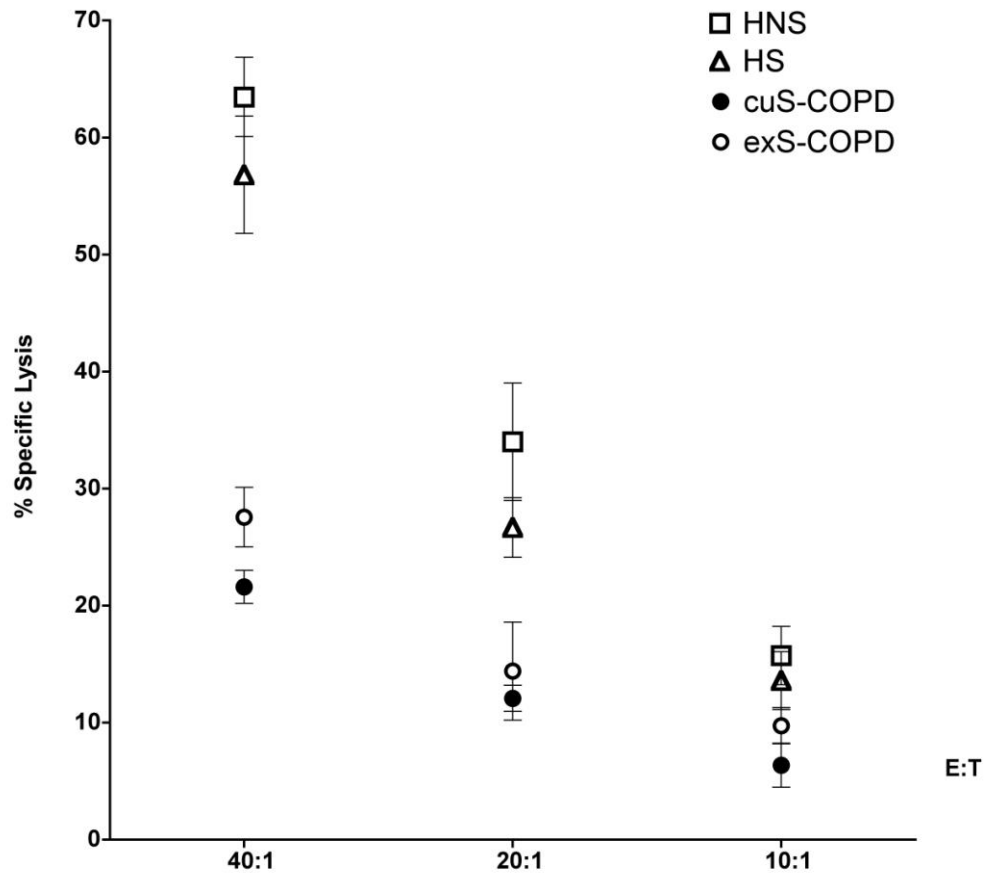


Figure 3.4.2 Cytotoxic activity of CD8⁺CD56⁺ T cells towards P815 cells in studied groups. CD8⁺CD56⁺ T cells were isolated from PBMCs of four interested groups, namely, health non-smokers (HNS, n=4), healthy smokers (HS, n=4), current smokers with COPD (cuS-COPD, n=3), and ex-smokers with COPD (exS-COPD, n=2). Purified CD8⁺CD56⁺ T cells were incubated with 2ug/ml of anti-CD3 for 24 hours before harvesting. Cytotoxicity was assessed after 4 hours at the indicated effector to target (E:T) ratio. Data are mean \pm standard deviation of different groups.

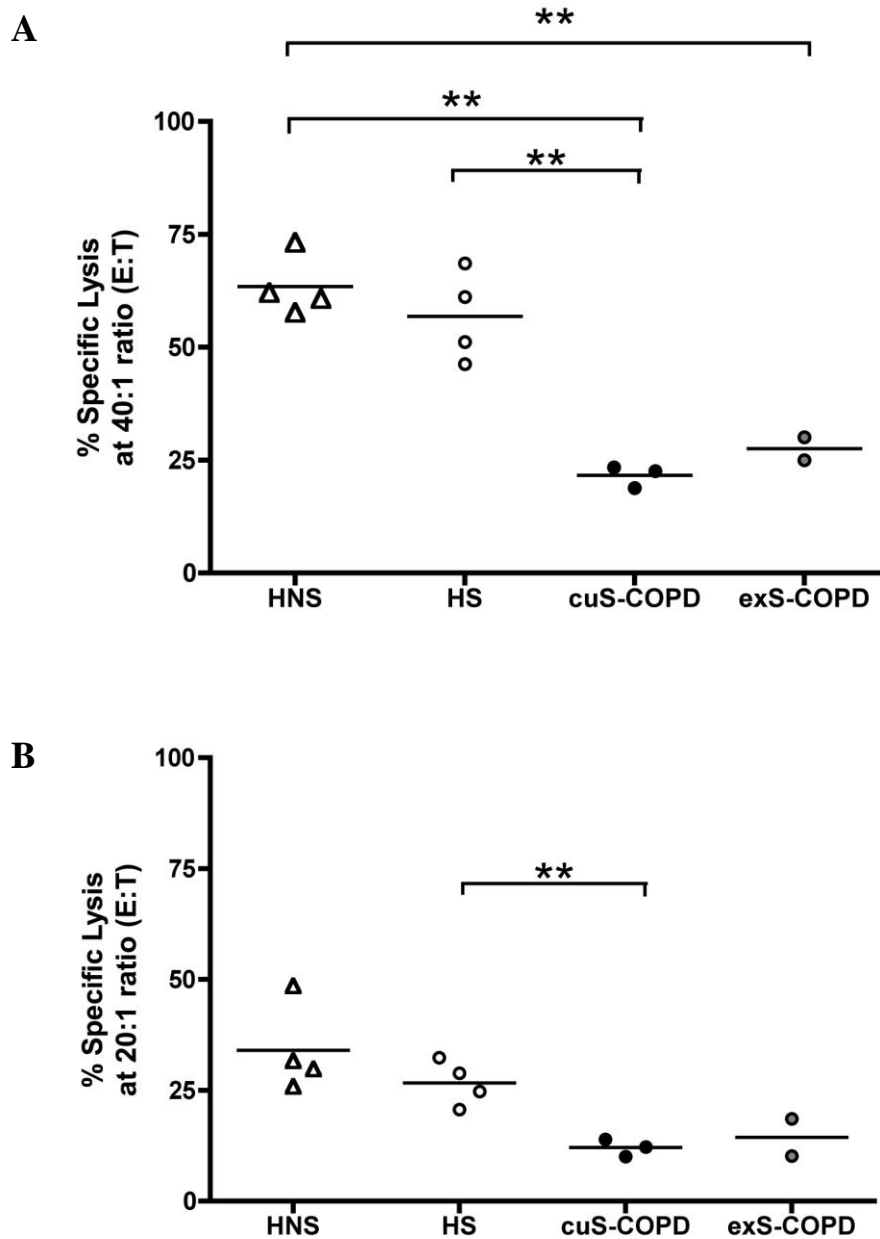


Figure 3.4.3 Cytotoxic activity of CD8⁺CD56⁻ T cells against P815 cells at the effector to target ratio of 40:1 (panel A) and 20:1 (panel B). Cytotoxicity detection for ratios 40:1 and 20:1 was performed in the same experiment in each sample. CD8⁺CD56⁻ T cells were immunomagnetically isolated from PBMCs of health non-smokers (HNS, n=4), current healthy smokers (HS, n=4), current smokers with COPD (cuS-COPD, n=3), and ex-smokers with COPD (exS-COPD, n=2). Significant differences were considered as p value<0.05 (*,p<0.01; **,p<0.005; ***,p<0.001).

Similar findings were also observed at ratio 20:1 (Figure 3.4.3.B). There were remarkable decreases on the cytotoxicity of CD8⁺CD56⁻ T cells from current smokers with COPD (12.1% \pm 1.9%) compared to those from healthy non-smokers (34.0% \pm 10.0%, $p < 0.01$) and current healthy smokers (26.7% \pm 5.1%, $p < 0.005$). There were no significant differences on the lysis ability of Tc cells between healthy non-smokers and current healthy smokers at each ratio.

3.4.4 Discussion

Here we demonstrated that CD8⁺CD56⁻ T lymphocytes from PBMCs showed significantly less cytotoxic activity against P815 cells in COPD patents compared to healthy non-smokers and healthy smokers. Also, dose dependent reductions were exhibited on the cytotoxicity of CD8⁺CD56⁻ T cells in each group. These findings were in accordance with an earlier study that showed the cytotoxicity of NK cells and NKT-like cells from PMBCs was significantly lower in COPD patients in contrast to healthy volunteers (Urbanowicz et al., 2009).

Target-cell death induced by CD8⁺CD56⁻ T cells can be initiated by various signals (Plataki et al., 2006). For example, when cytotoxic T cells are mixed with target cells and rapidly brought into contact by centrifugation, they can program

antigen-specific target cells to die by secretion of preformed effector molecules and cytokines, such as perforin, granzymes and INF- γ . K562 cell line was previously selected as the target cells in this research. However, no remarkable findings were observed between COPD patients and healthy groups. Previous studies have shown that K562 cells do not express HLA-A, B, C, or HLA-DR on their surface that are recognized by CD8⁺ T cells to induce cell cytotoxicity (Garson et al., 1985, Maziarz et al., 1990). Here we used P815 cell line as target cells. The lysis ability of CD8⁺ T cells against P815 cells was significantly reduced in COPD groups compared to healthy volunteers. This was supported by other studies that showed CD56⁺ T cells, which were NKG2D-positive, were not cytotoxic against K562 cells (Karimi et al., 2005, Trinchieri, 2003, Verneris et al., 2004). NKG2D might account for this reason. Besides MHC class I molecule dependant pathway, cytotoxic T cells also induce activating signal by MHC-unrestricted and TCR-independent pathway (Schmidt-Wolf et al., 1993). NKG2D is a representative MHC-unrestricted activation receptor expressed by human NK cells, CD8⁺ T cells and $\gamma\delta$ T cells. Multiple studies illustrate that NKG2D ligation leads to different functional outcomes on killer cells (Diefenbach et al., 2000, Groh et al., 2001, Jamieson et al., 2002, Pende et al., 2001). Groh *et al* showed that NKG2D augmented cytotoxicity

of CD8⁺ T cells only when cytotoxic T cells and target cells were MHC matched (Groh et al., 2001). Another research also confirmed that NKG2D-blocking antibodies dramatically attenuated the cytotoxicity of CD8⁺ T cells in tumor targets (Verneris et al., 2004).

DAP10 and DAP12 are two well-identified adapter proteins associated with NKG2D in human cytotoxic cells (Rosen et al., 2004). Studies confirm that NKG2D, together with the DAP10 and DAP12 molecules, has been shown to play a unique role in the mechanism of cytotoxicity of T cells (Karimi et al., 2005, Trinchieri, 2003, Verneris et al., 2004). Double knockdown of DAP10 and DAP12 reduced much of the killing ability of activated CD8⁺ T cells in PBMCs from healthy volunteers (Kakimi et al., 2000). However, DAP10 and DAP12 were associated with not only NKG2D receptor but also other receptors such as KIRs and CD94-NKG2C receptors (Lanier, 2009). DAP12-associated KIR has been identified on the subsets of human effector T cells in mice (Feng et al., 2005). Defining the DAP12-related KIRs in human are beyond our study, however, in future, the relationship between KIRs expression and cytotoxicity of killer cells may be another prospective study in COPD. Indeed, in an earlier section we showed that *ex vivo* expression of KIRs, but not NKG2D, on

CD8⁺ T cells were highly reduced in current smokers with COPD (Figure 3.1.8).

The relatively low participant numbers, especially the number of ex-smokers with COPD, should be pointed out. We did not find out any differences on the cytotoxic ability of CD8⁺CD56⁻ T cells between current smokers with COPD and ex-smokers with COPD. However, the significant differences between healthy and patient groups could partially explain the finding that the cytotoxicity of CD8⁺ CD56⁻ T cells was disease-related rather than smoking *per se*.

In summary, this study has shown that the cytotoxicity of CD8⁺CD56⁻ T cells was significantly reduced in the peripheral blood of COPD subjects (both current and ex) compared to healthy non-smokers and healthy current smokers.

3.5 Killer cell activation in bronchoalveolar lavage fluid from COPD

3.5.1 Introduction

In the last decade, there have been many studies performed on bronchial biopsies, lung parenchyma and peripheral blood obtained from COPD patients compared with healthy non-smokers and smokers with normal lung function. Bronchoscopy with bronchoalveolar lavage (BAL) is an important research tool for assessing airway inflammation in a variety of inflammatory lung diseases. Increased numbers of neutrophils (Battaglia et al., 2007), CD1a⁺ immature DCs (Casolaro et al., 1988, Soler et al., 1989), macrophages (Barnes, 2004d, Saetta et al., 1993, Saetta et al., 1997, Wallace et al., 1992), eosinophils (Saetta et al., 1996, Saetta et al., 1994), CD8⁺ T cells (Saetta et al., 2002, Saetta et al., 2003), γ/δ T cells (Ekberg-Jansson et al., 2000, Pons et al., 2005) and regulatory T cells (Smyth et al., 2007) were detected in the BAL fluid from either current healthy smokers or current smokers with COPD. However, the number of iNKT cells from BAL was significantly decreased (less than 2% of total CD3⁺ T cells) in subjects with stable or exacerbated COPD (Vijayanand et al., 2007). Besides immune cells,

abnormal levels of immune molecules were also detected in BAL from COPD patients. The concentration of CXCL1 (Traves et al., 2002), CXCL5 (Tanino et al., 2002), CCL2 (Capelli et al., 1999, de Boer et al., 2000, Traves et al., 2002), GM-CSF (Balbi et al., 1997), ET-1 (Bacakoglu et al., 2003, Chalmers et al., 1999), histamine (Kalenderian et al., 1988) and tryptase (Bessa et al., 2008, Kalenderian et al., 1988) were all significantly increased in BAL fluid from COPD patients. Levels of both IL-4 released by CD8⁺ T cells and IFN- γ released by CD4⁺ T cells were also elevated in the BAL fluid from COPD patients (Barczyk et al., 2006). Increased xanthine/xanthine oxidase activity has been observed in BAL fluid from COPD patients (Pinamonti et al., 1998). The expression of perforin and granzyme B on CD8⁺ T cells from BAL fluid were also significantly increased in COPD patients and healthy smokers compared non-smokers (Hodge et al., 2006).

To date, the study on the killer cell activation from BAL in COPD is rare. In this chapter, we aim to examine the activation levels of killer cells including CD8⁺ T cells, NKT-like cells and NK cells from BAL in COPD patients.

3.5.2 Material and Methods

3.5.2.1 Study population and procedures

The criteria of participant enrolments are the same as previously introduced. There were only four current smokers with COP included in this study. Table 3.5.1 shows the demographic and spirometric data of the participants.

3.5.2.2 BAL collection

Bronchoscopy with BAL was performed by a clinician from the Queen's Medical Centre. More details are provided in Section 2.8.

Table 3.5.1 Demographic and spirometric values of the BAL study

Current smokers with COPD	Age (years)	Gender (M/F)	Packs/year	Chronic bronchitis (Yes/No)
n=4	49 (43-57)	2/2	41 (38-46)	4/0

FEV1 (% pred)	FEV₁/FVC (%)	ΔFEV₁ post bronch	BMI (kg/m²)
61 (50-72)	37 (31-44)	1.6 (1.4-1.9)	19.7 (16.6-22.5)

3.5.2.3 BAL cell collection

Before centrifugation of BAL fluid, two 50ml centrifuge tubes were weighted on an electronic scale. Samples were pipetted into the tubes and volumes of them were noted. Filled tubes were reweighted and the weight of recovered BAL fluid was calculated. More details are provided in Section 2.8.

3.5.2.4 Extracellular staining

To determine the cell populations in the BAL, extracellular staining for these cells were performed as described previously. Cells were labeled with fluorescently conjugated antibodies including CD3 (PC7), CD8 (ECD) and CD56 (PE) to identify the proportion of cell population of interest. CD25 (FITC) and CD69 (PC5) were also added to detect the activation of interested cell types. All samples were analysed by flow cytometry.

3.5.2.5 Flow cytometric analysis

Flow cytometric analysis of antibody labelled cells was performed using an EPICS Altra (Beckman Coulter, Luton, UK). More details are provided in Section 2.9.

3.5.2.6 Statistical analysis

The statistical analysis was performed with Prism software, version 4.0c (GraphPad). Differences between four cell types including CD8⁺ T cells, CD4⁺ T cells, NKT cells and NK cells from COPD smokers were tested using the parametric student's *t* test. Confidence interval 99% was used to show two-tailed *p* values. *P* values of less than 0.01 were considered to indicate statistical significance.

3.5.3 Results

Killer cell activation in BAL from COPD patients ex vivo

All individuals had similar total mononuclear cell numbers, which were within the normal range (data not shown). Representative flow analysis dot plots showed Forward Scatter (FS) vs Side Scatter (SS) for identification of live and dead cells (Figure 3.5.1.A), along with dot plots for identification of CD8⁺ T-lymphocytes (CD3⁺CD8⁺), CD4⁺ T-lymphocytes (CD3⁺CD4⁺), NKT-like cells (CD3⁺CD56⁺), and NK cells (CD3⁻CD56⁺) (Figure 3.5.1.B). In addition dot plots for CD69 and CD25 staining of these four cell types from BAL of a COPD patient are presented in Figure 3.5.2. Cells are designated activated when they express one or both of the activation markers (i.e., CD69 positive, CD69/CD25 double positive or CD25 positive). CD8⁺ T cells (92.0%, 89.1-97.1; *p*<0.001),

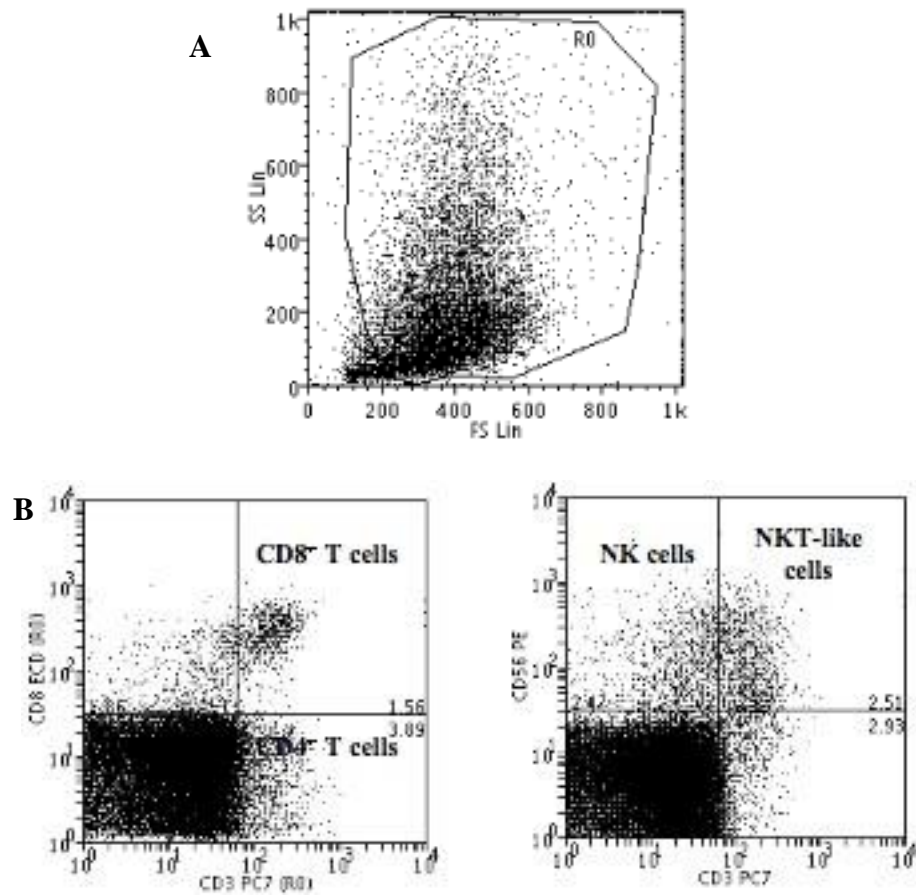


Figure 3.5.1 Representative dot plots gating for live/dead cells and identification cell populations from current smokers with COPD. (A) Forward Scatter (FS) vs Side Scatter (SS) plot for identification of live and dead cells; (B) Identification of CD8⁺ T cells (CD3⁺CD8⁺), CD4⁺ T cells (CD3⁺CD4⁺), NKT-like cells (CD3⁺CD56⁺) and NK cells (CD3⁻CD56⁺).

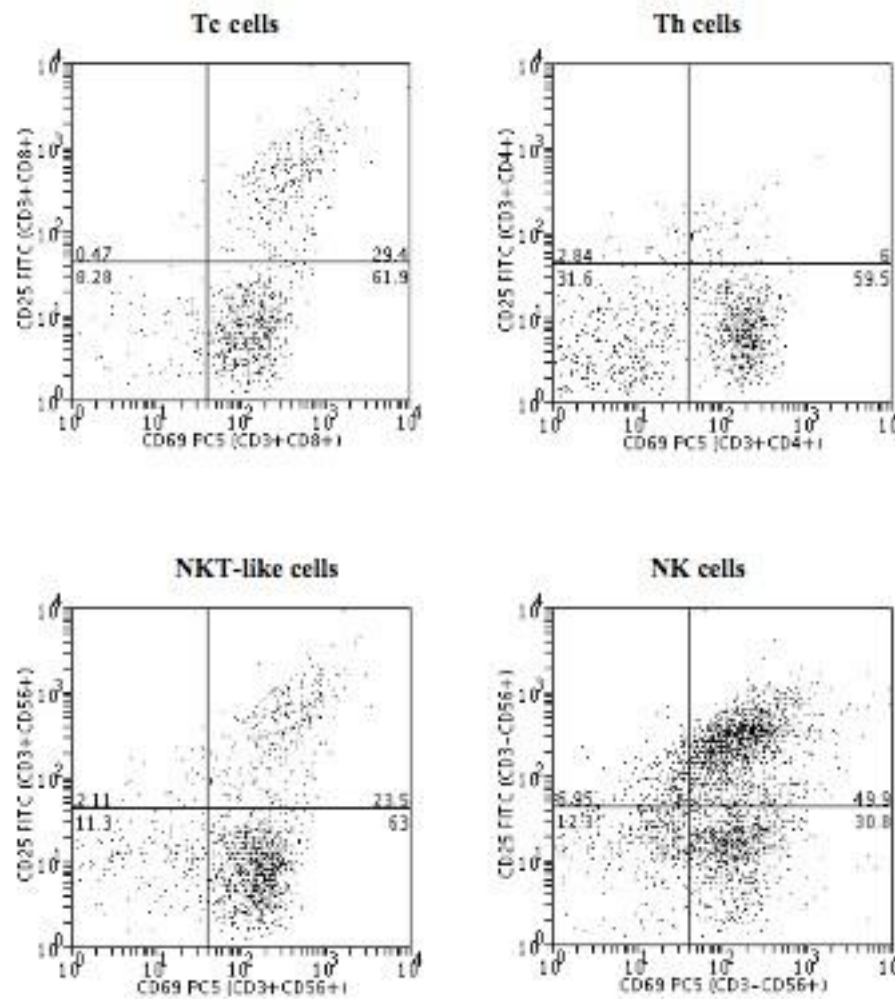


Figure 3.5.2 Representative dot plots for identification of activation of cells of interest from a current smoker with COPD. Activation is designated by expressing either CD69 alone (lower right quadrant), CD69 and CD25 (upper right quadrant) or CD25 alone (upper left quadrant).

NKT-like cells (96.5%, 87.7-99.2; $p<0.001$) and NK cells (94.5%, 88.7-97.3; $p<0.001$) were all highly and significantly activated in compared to $CD4^+$ T cells (46.9%, 40.2-68.5) in COPD patients (Figure 3.5.3). CD69 and CD25 represent early and late activation markers for cell activation respectively. As shown in Figure 3.5.4, activated $CD8^+$ T cells and NK cells primarily express CD69, whereas the major proportion of activated NKT-like cells is CD69 and CD25 double positive. All cells analysed expressed low level of CD25.

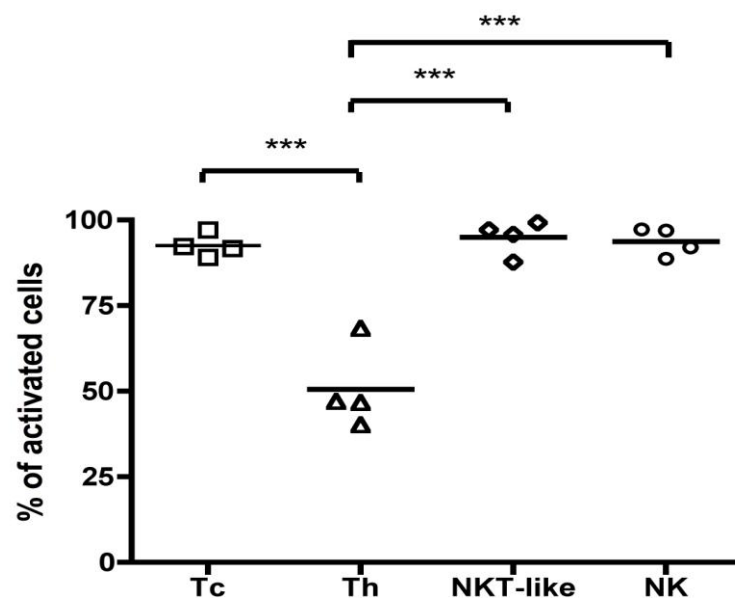


Figure 3.5.3 Activation of cells of interest including $CD8^+$ T cells (Tc), $CD4^+$ T cells (Th), NKT-like cells and NK cells from BAL fluid of current smokers with COPD (n=4). Tc, NKT-like and NK cells were all significantly activated compared Th cells from BAL of current smokers with COPD ($p<0.001$).

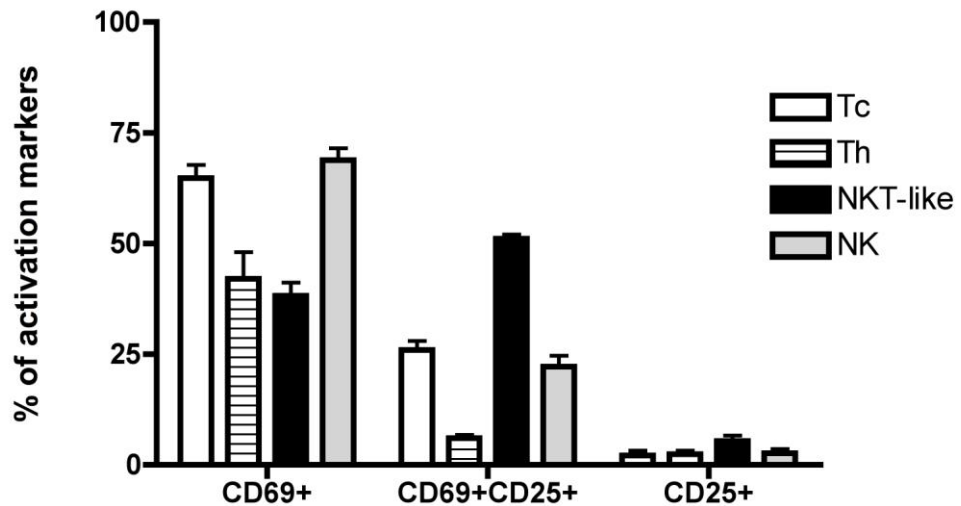


Figure 3.5.4 Activation stages of cells of interest including CD8⁺ T cells (Tc), CD4⁺ T cells (Th), NKT-like cells and NK cells from BAL fluid of current smokers with COPD (n=4). Tc, NKT-like and NK cells were all significantly activated compared Th cells from BAL of current smokers with COPD.

3.5.4 Discussion

Bronchoscopy with BAL is an important research tool for assessing airway inflammation in a variety of inflammatory lung diseases. Many studies have shown the abnormal numbers of immune cells and molecules from BAL in COPD. Here we showed that killer cells including CD8⁺ T cells, NKT-like cells and NK cells were all highly activated compared to CD4⁺ T cells from BAL of current smokers with COPD. This finding was supported by other studies. CD8⁺ T-lymphocytes from BAL fluid of smokers with or without COPD were significantly activated compared to healthy non-smoking

volunteers (Roos-Engstrand et al., 2009). Also, increased proportion of CD8⁺ T cells expressing CD69 was observed in the lung tissue of COPD patients compared to non-smokers (Freeman et al., 2010). However, in our study we could not conclude that high levels of killer cell activation were disease specific due to lack of healthy controls.

CD69 and CD25 represent early and late activation markers respectively for cell activation. Interestingly, we found that the major proportion of activated NKT-like cells was CD69 and CD25 double positive in BAL of current smokers with COPD, whereas CD8⁺ T cells and NK cells mainly express CD69 (early activation marker) *ex vivo*. This finding was partially supported by Freeman *et al* study. They found increased proportion of CD8⁺ T cells expressing CD69 in the lung tissue of COPD patients compared to non-smokers (Freeman et al., 2010). NKT cells, a subset of T lymphocytes but also express NK cell surface markers, exhibit both pro-inflammatory and immuregulatory characteristics. Cytokines released by NKT cells are always rapid and transient, with a peak of activity within 24 hours. After 24 hours, productions of these cytokines, for example IFN- γ , quickly return to baseline levels (Eberl and MacDonald, 1998). This may explain why NKT cells were more quickly activated than CD8⁺ T cells and NK cells.

Here we also showed the activation level of CD4⁺ T cells was lower compared to interested killer cells in COPD patients. This is consistent with our findings in PBMCs and sputum that both showed low level activation of Th cells in both healthy groups and COPD subjects. However, there were no control groups in BAL study. In future, we could compare the expression of activation markers of CD69 and CD25 between healthy volunteers and patient groups. Also, to identify whether those CD4⁺ T cells were regulatory T cells, FoxP3 staining could be applied.

Overactivation of killer cells can induce hypersecretion of inflammatory cytokines and chemokines that consequently cause other types of cell activation, and finally contribute to the lung tissue damage in the COPD patients. Although here we could not conclude these highly activated cells were disease related because of no control samples, in future, it may be worth investigating the killer cells activation and their functional receptor expression in BAL fluid from both healthy volunteers and COPD patients, which provide new insights on the role of killer cells and contributing to a better knowledge of the pathogenesis of COPD.

Discussion

COPD is characterised by poorly reversible airflow obstruction and airway inflammation with high morbidity and mortality (Celli and MacNee, 2004). As an inflammatory disease, both innate and adaptive immune system show abnormal features in COPD (O'Shaughnessy et al., 1997, Saetta et al., 1998, Baraldo et al., 2004, Barnes, 2004b, Demedts et al., 2007, Urbanowicz et al., 2009) that eventually lead to chronic obstructive bronchiolitis, emphysema and mucus hypersecretion.

Activated neutrophils (Damiano et al., 1986, Galdston et al., 1977), macrophages (Molet et al., 2005, Montano et al., 2004), DCs (Demedts et al., 2007) and CD8⁺ T lymphocytes (Roos-Engstrand et al., 2009) have been reported in airways or sputum of COPD patients compared to normal non-smokers. However, studies on the activation levels of killer cells including CD8⁺ T cells, NKT-like cells and NK cells from PBMCs are unexplored. Previous studies also demonstrated the potential involvement of the three main classes of human killer cells, namely, CD8⁺ T-lymphocytes, NKT-like cells and NK cells in COPD (Paats et al., 2012, Roos-Engstrand et al., 2009, Roos-Engstrand et al., 2010, Urbanowicz et al., 2009) that play key roles in inflammatory responses.

Here we showed, *ex vivo*, for the first time that activation phenotype of these cells in the peripheral blood. Activation levels CD8⁺ T-lymphocytes, NKT-like cells and NK cells were significantly higher in current smokers with or without COPD in comparison with healthy non-smokers and ex-smokers with COPD. This finding indicated a systemic influence of smoking on killer cell activation, irrespective of the presence of airflow limitation. We also found that NK cell activation was positively correlated with the number of cigarettes currently smoked in COPD patients. No correlations between cell activation and lung function (FEV₁ percentage of predicted) were found. These data indicate that in peripheral blood, cell activation levels are more related to current smoking habit rather than disease. A previous study examining the effect of smoking cessation has shown that CD8⁺ T-lymphocytes from BAL fluid of smokers with or without COPD expressed high levels of CD69 and CD25 activation markers (Roos-Engstrand et al., 2009). Also, an increased proportion of CD8⁺ T cells expressing CD69 was observed in the lung tissue of COPD patients compared to non-smokers (Freeman et al., 2010). Sputum T-lymphocytes have been shown to express higher levels of CD103 and CD69 than blood lymphocytes suggesting higher numbers of activated intraepithelial phenotype T-lymphocytes in the lung of COPD patients (Leckie et al., 2003).

Also, in our study, CD4⁺ T cells expressed very low levels of activation markers, namely, CD69 and CD25, in all groups. CD25 is also a cell marker for regulatory T cells. Therefore, we hypothesised those CD4⁺CD25⁺ T cells might mainly be regulatory cells rather than activated cells. To examine this further we generated Forward Scatter (FS) vs Side Scatter (SS) plots that showed, unlike other activated cell types expressing CD25 that were larger and more granular, those CD4⁺CD25⁺ T cells were primarily small and less granular and thus quite probably inactivated lymphocytes. However, to confirm this observation further studies are required, such as staining for FoxP3. Although two studies have found highly activated Th1 cells in the lung of ex-smokers with COPD and in the small airway wall of smokers with severe COPD by detecting their chemokine and cytokine secretion, those samples are actually from COPD patients with emphysema or co-suffering from cancer (Turato et al., 2002).

KIR (CD158e1) expression by killer cells was significantly lower in peripheral blood of current smokers with or without COPD compared to healthy non-smokers. This finding was opposite to the activation levels of these killer cells. Signalling balance between activating receptors and inhibitory receptors is one of the crucial factors determining cell activation. Similar

finding was shown in Hodge *et al* study. CD94 is another inhibitory receptor expressed by NKT-like cells and NK cells (Lazetic *et al.*, 1996). Decreased expression of CD94 on both NKT-like cells and NK cells were detected in PBMCs from COPD patients compared to healthy volunteers (Hodge *et al.*, 2012). However, they did not detect any significant differences on the cell activation levels of NKT-like and NK cells (only CD69 expression) between groups. Therefore, our findings provided the first demonstration of association of cell activation and KIR expression.

We also showed that activating receptor NKG2D was highly expressed by CD8⁺ T cells, NKT-like cells and NK cells but not CD4⁺ T cells, which was in accordance with previous studies that demonstrate NKG2D receptors are constitutively expressed and almost exclusively on cytotoxic cells including CD8⁺ T cells, NKT cells, NK cells and T cells (Maasho *et al.*, 2005). There were no significant differences of NKG2D expression on CD8⁺ T cells, CD4⁺ T cells, NKT-like cells and NK cells respectively, in the four groups. This further supports the proposal of increased peripheral blood killer cell activity in smokers due to the increased ratio of activating to inhibitory receptors.

Of note, our group has published data showing, *ex vivo*, the activation levels of killer cells in sputum samples from COPD patients are elevated (Wang et al., 2013). Unlike systemic activation, lung activation is due specifically to COPD, irrespective of current smoking status.

In this study, activation levels of cells from BAL of COPD patients were examined. CD8⁺ T cells, NKT-like cells and NK cells were all highly activated compared to CD4⁺ T cells in BAL fluid from COPD patients *ex vivo*. This finding was similar to those we observed in PBMCs from current smokers with COPD. Moreover, CD8⁺ T cells and NK cells showed early activation stage (mainly expressing CD69), whereas the main proportion of activated NKT-like cells were CD69 and CD25 double positive. However, it was hard to conclude whether it was COPD specific due to no control samples from healthy participants. This is because the procedure of taking BAL samples is indeed very uncomfortable for those volunteers.

Cytotoxicity of CD8⁺ T cells was also investigated in our study. We demonstrated that CD8⁺CD56⁻ T lymphocytes from PBMCs showed significantly less cytotoxic activity against P815 cells in COPD patients compared to healthy non-smokers and healthy smokers. Also, dose dependent reductions were

exhibited on the cytotoxicity of CD8⁺CD56⁻ T cells in each group. This finding was in accordance with previous research on lysis ability of CD56⁺ cells. The cytotoxicity of CD56⁺ cells, including NK cells and NKT-like cells, from PMBCs against K562 cell line was significantly lower in COPD patients in contrast to healthy volunteers (Urbanowicz et al., 2009).

Of note, K562 cell line was previously selected as target cells in this research. However, no remarkable findings were observed between COPD patients and health groups. Here we used P815 cell line as target cells. Interestingly, the lysis ability of CD8⁺ T cells against P815 cells was significantly reduced in COPD groups including both current and ex-smokers with COPD compared to healthy volunteers. Anti-CD3 monoclonal antibody induces cytolysis by cross-linking CD3⁺ effector and IgG-FcR⁺ target cells. After coating with P815 cells, anti-CD3 can induce P815 cells bearing the relevant IgG-FcR and further perform as antigen presenting cells to activate CD8⁺ T cells (van de Griend et al., 1987). Furthermore, other studies that showed NKG2D-positive CD56⁻ T cells were not cytotoxic against K562 cells (Karimi et al., 2005, Trinchieri, 2003, Verneris et al., 2004). Besides MHC class I molecule dependant pathway, cytotoxic T cells also induce activating signal by MHC-unrestricted and TCR-independent pathway

(Schmidt-Wolf et al., 1993). Groh *et al* showed that NKG2D augmented cytotoxicity of CD8⁺ T cells only when cytotoxic T cells and target cells were MHC matched (Groh et al., 2001). Another research also confirmed that NKG2D-blocking antibodies dramatically attenuated the cytolysis of CD8⁺ T cells in tumor targets (Verneris et al., 2004).

We did not find any differences on the cytotoxic ability of CD8⁺CD56⁻ T cells between current smokers with COPD and ex-smokers with COPD. However, the significant differences between healthy and patient groups could partially reveal the finding that the cytotoxic of CD8⁺ CD56⁻ T cells was disease-related rather than smoking *per se*.

Cigarette smoking, as the major risk factor in the development of COPD, affects structural and functional changes in the respiratory ciliary epithelium (Cantin et al., 2006, Mio et al., 1997, Wyatt et al., 1999), lung surfactant protein (Honda et al., 1996, Subramaniam et al., 1996) and immune cells (Mohan, 2002a).

Here we investigate the impact of cigarette smoke extract (CSE) on α -CD3, IL-2 or IL-15 activated killer cells including Tc cells, NKT-like cells and NK cells, and further compared the

absolute suppression levels induced by CSE on those cells upon different activation between four groups, namely, healthy non-smokers, current healthy smokers, current smokers with COPD and ex-smokers with COPD.

Optimal concentration of each activator and incubation time for cell culture was achieved by a preliminary protocol. Tc cells, NKT-like cells and NK cells were all highly activated when the concentration of α -CD3 increased to 1ug/ml. However, in preliminary experiments, dose-dependent increases in the activation levels of NK cells were also observed, although NK cells do not express CD3/TCR complex on their cell surface. To our knowledge, this could be easily explained. α -CD3 has the capacity of activating CD4⁺ T cells, Tc cells and NKT cells that further release immune mediators such as IL-2 or IL-15 (Smyth et al., 2005, Thompson et al., 1989). Furthermore, CSE also has effects on DCs and macrophages that also secrete high amount of IL-15 (Carson et al., 1995, Jonuleit et al., 1997). Both IL-2 and IL-15 are good sources for NK cell activation (Waldmann, 2006).

In our study, activation levels of killer cells were not significantly different in the absence or presence of CSE at increasing dilutions in healthy volunteers, without other

stimulations. These findings were in agreement with other groups who detected the CSE effects on DCs or NK cells from healthy volunteers (Mian et al., 2009, Mortaz et al., 2009). According to previous studies and our detection, we selected 1% of CSE for further experiments because it did not cause any toxic effects as revealed by MMT assay (Mian et al., 2008, Mian et al., 2009, Stringer et al., 2007).

The effects of CSE on killer cells were explored not only in healthy volunteers but also COPD patients in our study. Three significant results should be highlighted. Firstly, CSE markedly suppressed activation levels of IL-15 treated cells for all cell types in all groups. Secondly, in contrast to Tc cells and NKT cells, NK cells were the only cell type which activation levels upon different stimulation were all significantly decreased when CSE was added in all groups. Thirdly, current smokers with COPD were the only group that showed CSE had statistically significant effects on all interested cell types activation upon different stimulation. All above research were carried out in whole PBMCs. These findings were partially supported by different studies. Mian and colleague have shown that in PBMCs from healthy non-smokers, the expression of CD69, a key activation marker for NK cells, was significantly inhibited when PBMCs was treated by cigarette

smoke after poly I:C activation (Mian et al., 2009). Hogan *et al* demonstrated that CSE significantly reduced the cytokine production and the ability to kill target cells of α -GalCel activated invariant NKT cells in PBMCs from healthy non-smoking participants. Meanwhile, the activation marker of CD107, but not CD69, was decreased (Hogan et al., 2011).

CSE suppression effects on isolated NKT-like cells ($CD3^+CD56^+$) and NK cells ($CD3^-CD56^+$) in response to IL-15 were also compared between groups. We found that, *in vitro*, CSE significantly suppresses the activation levels of IL-15 treated NKT-like cells and NK cells isolated from PBMCs in healthy non-smokers and smokers with or without COPD. However, this effect on NK cells was much more significant in current smokers with COPD. Previous study has shown that cigarette smoke inhibited IL-12 and type-1 IFN production in activated PBMCs, which consequently decreased the IL-15 levels in healthy non-smokers (Mian et al., 2008, Mian et al., 2009). Also, both *in vitro* and *in vivo*, cigarette smoke significantly diminished IL-15 mRNA levels in poly I:C-activated PBMCs from healthy non-smoking participants. IL-15 production was markedly impaired in smoke-conditioned medium treated PBMCs upon poly I:C stimulation.

Granzymes, perforin and Fas-Fas ligand are three major killing molecules that facilitate NKT-like cells and NK cells performing their cytotoxic functions. We found that expression of granzyme B on IL-15 treated NKT-like cells and NK cells were markedly reduced by CSE in PBMCs from healthy non-smokers. Also, CSE significantly suppressed granzyme B expression by NK cells in response to IL-15 in current smokers with COPD. Surprisingly, CSE did not induce any remarkable impacts on expression levels of perforin and Fas-ligand in all groups. Perforin was highly expressed in all stimulating conditions whereas Fas-L showed very low levels. These findings were partially supported by other research. Morissette and colleague illustrated that no statistical differences were observed on the expression of perforin and granzyme B in peripheral NK cells from either healthy volunteers or smokers with evidence of emphysema. Protein expression of Fas-ligand was not detected by flow cytometry (Morissette et al., 2007). The cytotoxicities of NK cells and NKT cells from PBMCs were both markedly reduced in COPD patients compared to healthy volunteers (Urbanowicz et al., 2009). *In vitro*, the percentage of perforin /granzyme B double positive NK cells was significantly lower in COPD patients compared to controls. Although there was a markedly increase in soluble Fas in plasma from severe COPD patients, there were no differences

on its expression in mild or moderate COPD and healthy controls (Yasuda et al., 1998). Hodge *et al* also demonstrated that there were no significant changes in the expression levels of granzyme A by NK or NKT-like cells in the blood of COPD patients compared to controls (Hodge et al., 2013). Taken together, our data suggest that granzyme B, but not perforin and Fas-ligand, was a functional molecule attributing to the significant impairment of NK cell killing ability in response to CSE in COPD patients. The action of granzyme B against target cells can be either perforin dependent (Bossi and Griffiths, 2005, Monks et al., 1998) or perforin-independent (Froelich et al., 1996, Metkar et al., 2002, Shi et al., 1997). No matter which action is undertaken, granzyme B mediated apoptosis is caspase-dependent (Talanian et al., 1997, Yang et al., 1998).

We also showed, *in vitro*, there were no significant differences on KIR (CD158e1) expression by CD56⁺ with or without CSE treatment. KIR expression was at a very low level in each group. Few studies were carried out to detect the KIRs expression in either healthy volunteers or COPD patients. However, one study reported that NKG2D expression was not inhibited in cigarette smoke treated PBMCs after poly I:C induction (Mian et al., 2009). Therefore, KIR (CD158e1) was

not a functional inhibitory receptor for CSE induced suppression of CD56⁺ cells activation in response to IL-15.

Although CSE significantly suppressed the isolated CD56⁺ cell activation and their granzyme B expression in IL-15 conditioned medium in both healthy non-smokers and current smokers with COPD, there were no statistical differences on the absolute suppression levels between groups, which were different from our early findings that showed in un-isolated NK CD56⁺ cells, absolute suppression levels induced by CSE on IL-15 mediated NK cell activation was significantly higher in current smokers with COPD compared to healthy non-smokers and current healthy smokers. Two reasons might be accounted for that. Firstly, small sample size was used in this study. In figure 3.3.7, the absolute suppression level induced by CSE on NK cell activation was higher than other two healthy groups, although no statistical differences were presented. Secondly, IL-15 is a potent cytokine with the ability of stimulating many kinds of immune cells, such as memory CD4⁺ T cells, naive or memory CD8⁺ T cells, DCs, NKT cells and NK cells, which can further secrete more cytokines and chemokines to induce adaptive immune responses (Anderson et al., 1995, Cosman et al., 1995). Therefore, in whole PBMCs

sample, IL-15 can further induce other immune response on un-isolated CD56⁺ cells.

It is important to note that cigarette smoke comprises a complex mixture of more than 4000 chemicals, such as nicotine, carbon monoxide, and tar (Hoffmann and Wynder, 1986, Stedman, 1968). Their immune-modulation effects on immune cells are distinct and also dose-dependent (Holt et al., 1976). To determine the specific component of cigarette smoke that regulates killer cell activation upon different stimulation is beyond the scope of our studies.

Several limitations should be considered in the whole project. Firstly, in the *ex vivo* study, there are many kinds of inhibitory and activating receptors expressed on killer cells but due to limited cell numbers we analyse two representative receptors that give an indication of the killing function of these cells. Secondly, we could not obtain healthy samples in our BAL study. Thirdly, *in vitro*, only small numbers of subjects were recruited. Moreover, there are many cytotoxic granules such as granzyme A and granzyme K related to killing mechanisms of CD56⁺ cells. However, proportions of NKT-like cells and NK cells are very low in PBMCs. We could just detect

representative killing molecules in our research due to cell numbers.

In conclusion, five important findings have been illustrated in this study. Firstly, we showed for the first time that, *ex vivo*, killer cells, including CD8⁺ T cells, NKT-like cells and NK cells, were significantly activated in current smokers with or without COPD compared to healthy non-smokers. Secondly, KIR (CD158e1) expression by all killer cell types was dramatically lower in smokers with or without COPD in comparison with healthy non-smokers. Thirdly, the cytotoxicity of CD8⁺ T cells from COPD patients were significantly less than that in healthy volunteers. Forthly, *in vitro*, CSE markedly decreased IL-15 treated NK cell activation in current smokers with COPD compared to other three groups. Finally, the expression of granzyme B, but not perforin and Fas-L, was also significantly inhibited on IL-15 stimulated NK cells when CSE was added.

Over activation of killer cells provide more understanding on the pathology of COPD development. Abnormal expression of KIR also suggests the potential use of KIRs therapy in COPD. Less cytotoxicity of CD8⁺ T cells may explain why lung tissue is susceptible to infection in COPD patients. Suppression effects induced by CSE on NK cell activation and killing

molecule expression indicate that chronic exposure to cigarette smoke leads to not only cellular death and damage in the airways, but also failure to activate critical intracellular signalling mechanisms that initiate protective host defence. The consequence is dysfunction of various cell types with accompanying predisposition to respiratory infection.

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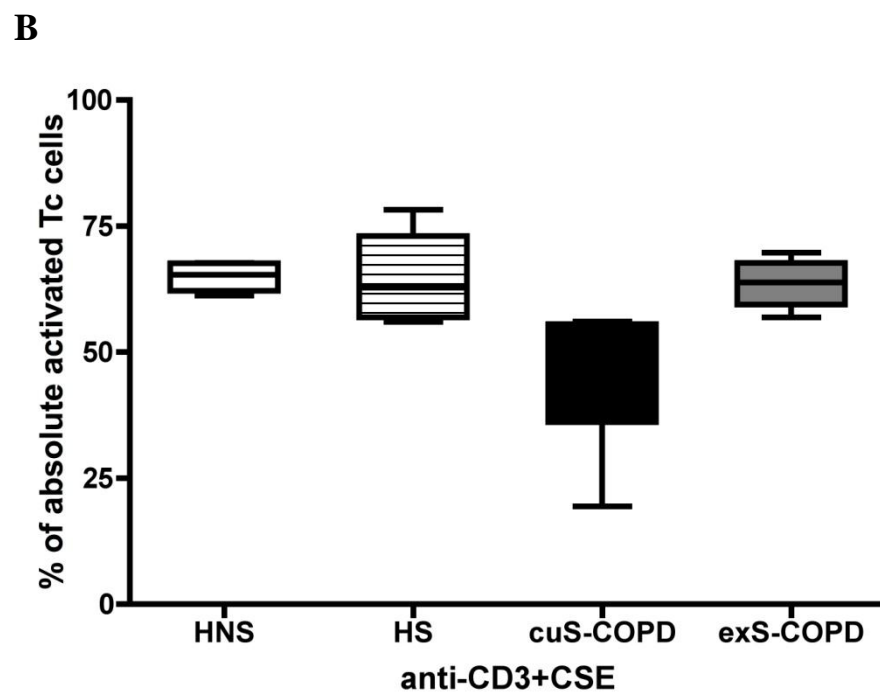
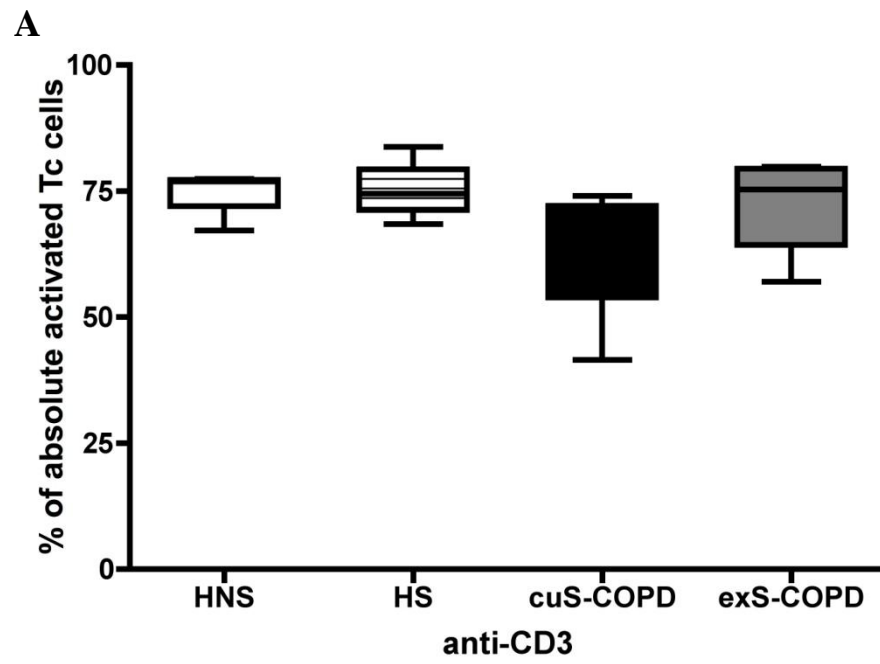
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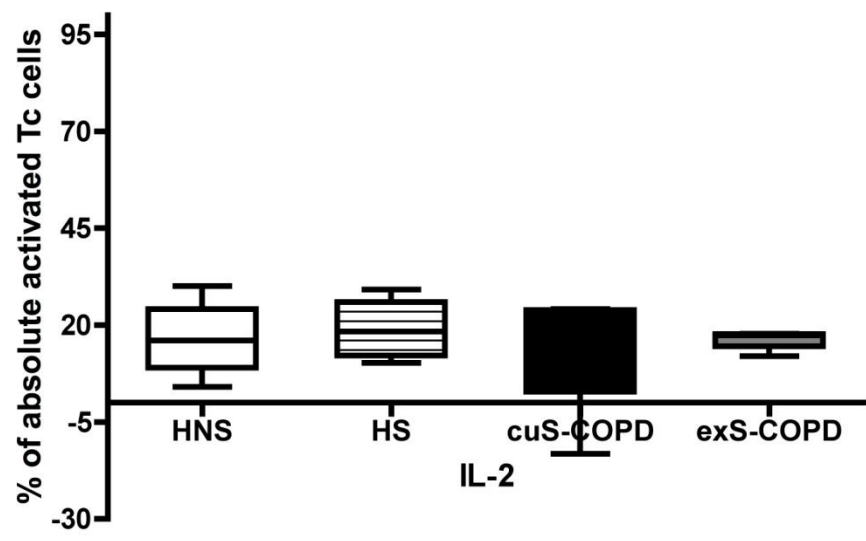
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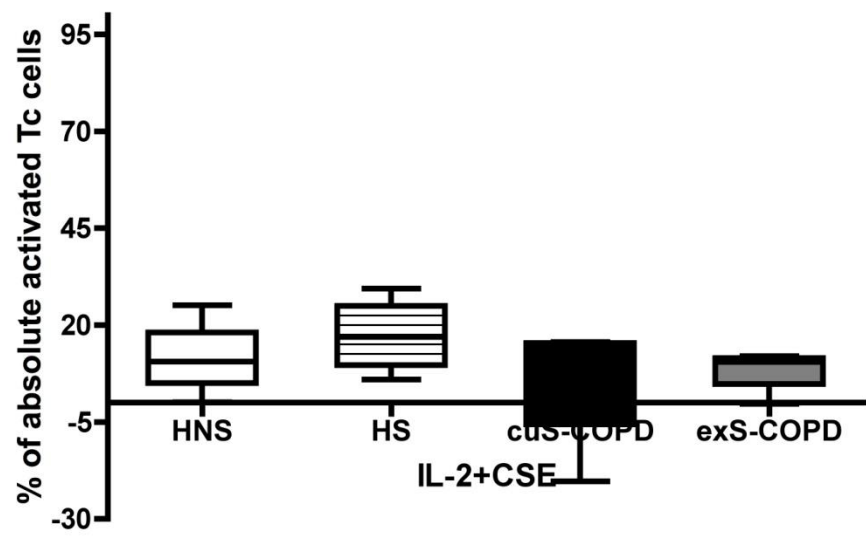
Appendix



C



D



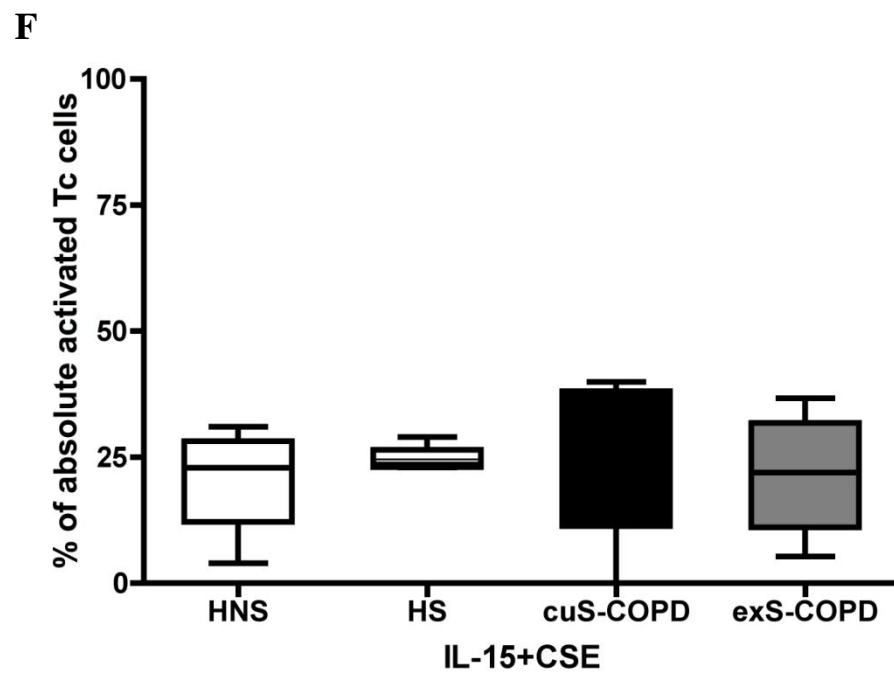
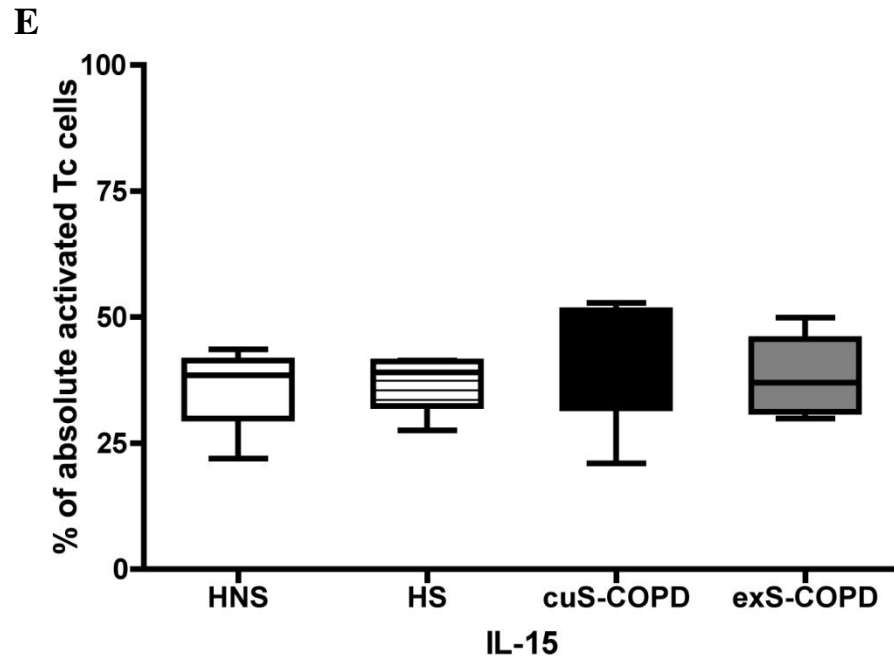
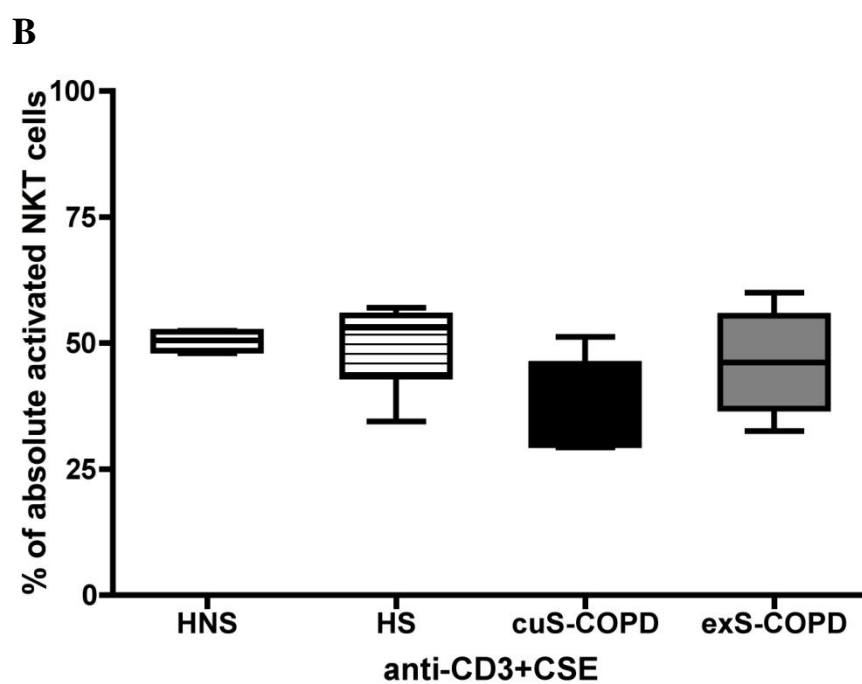
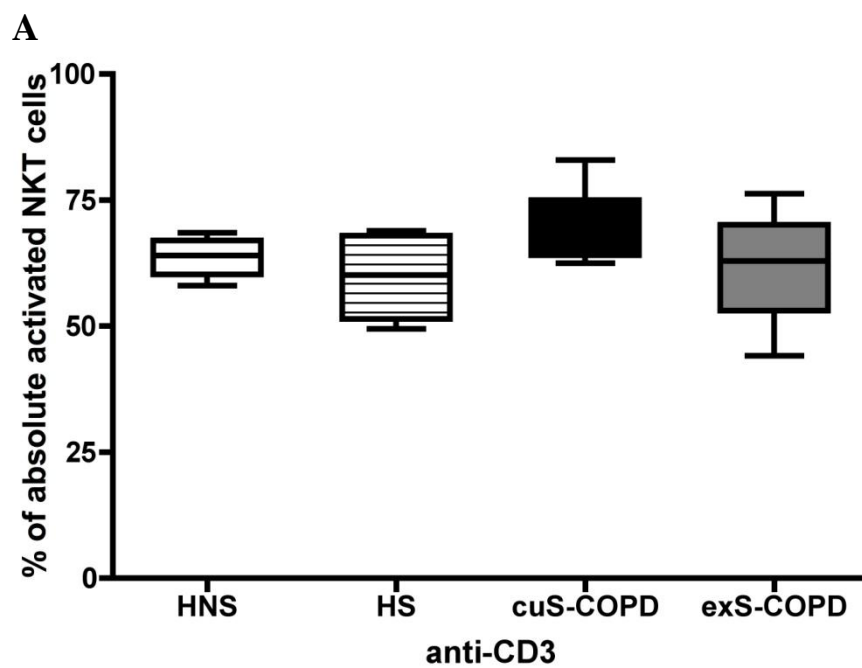
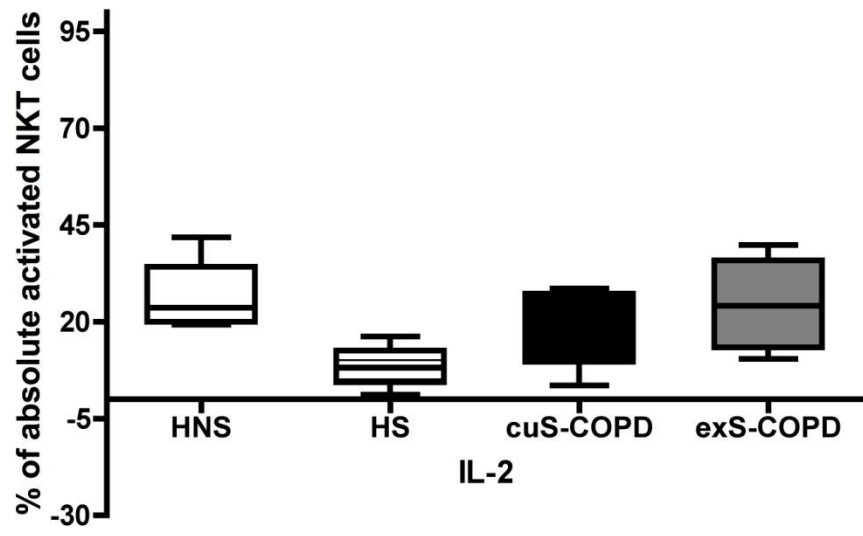


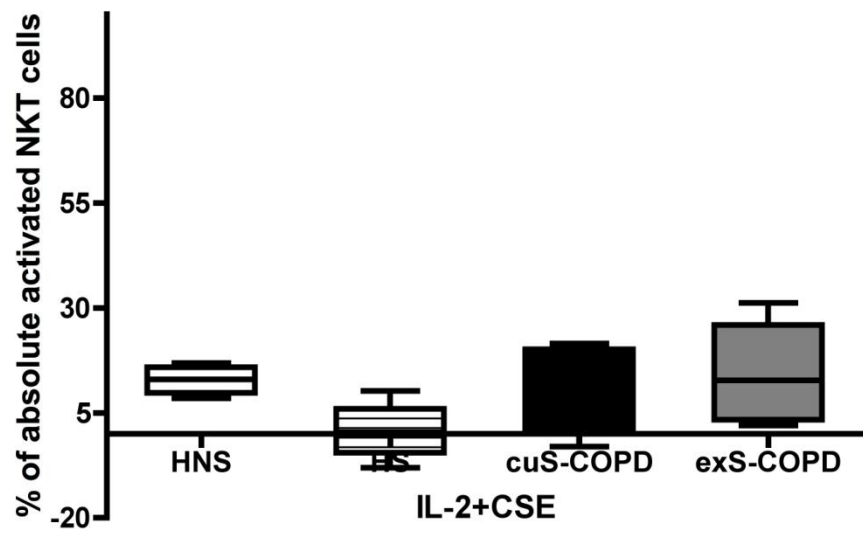
Figure 1. Comparison of absolute activation levels of CD8⁺ T cells upon different stimulation in four groups including healthy non-smokers (HNS, n=4), current healthy smokers (HS, n=4), current smokers with COPD (cuS-COPD, n=4) and ex-smokers with COPD (exS-COPD, n=4). PBMCs were collected and incubated with different reagents for 48 hours.



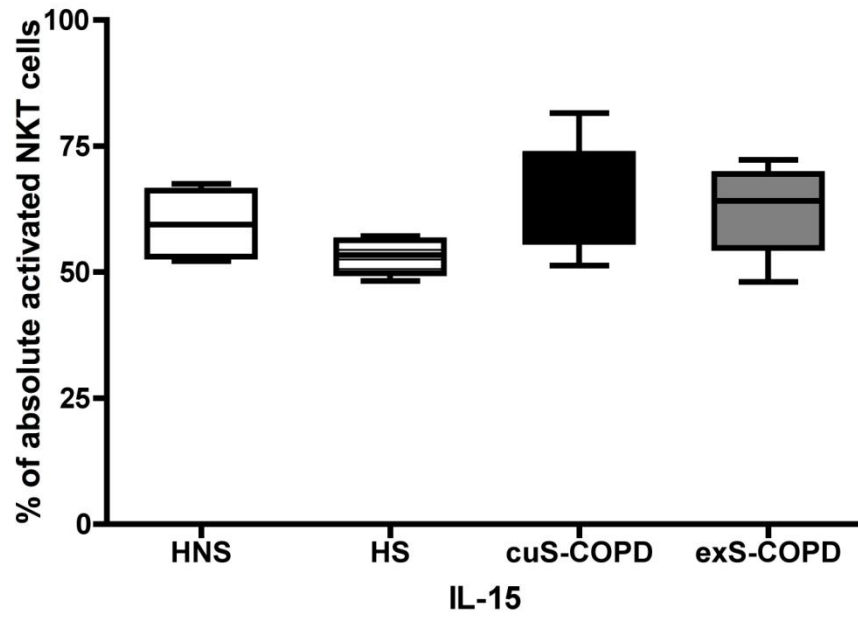
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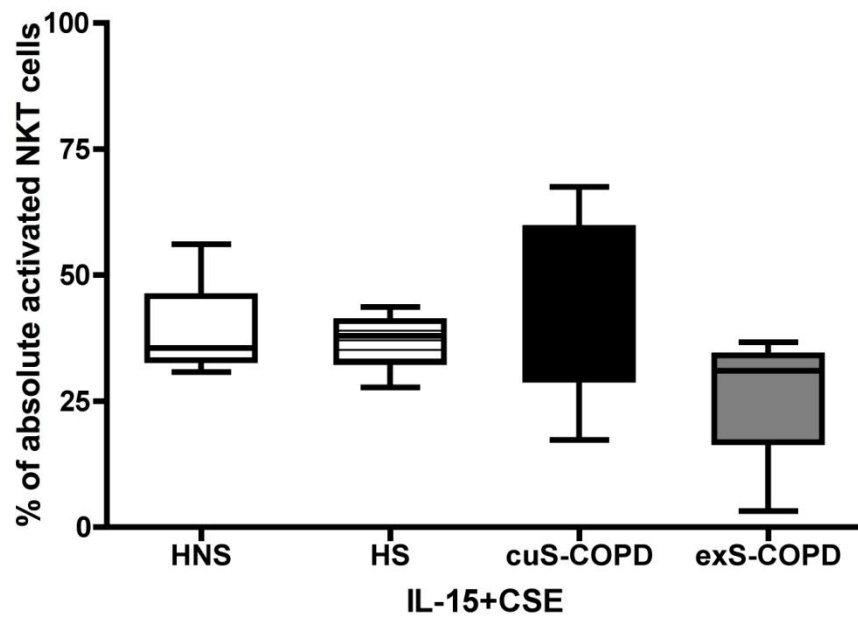
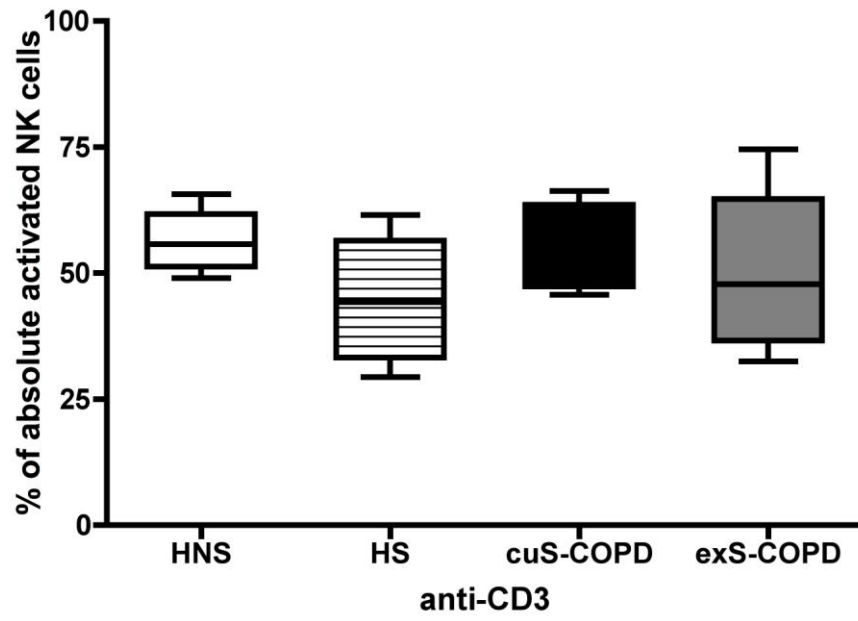
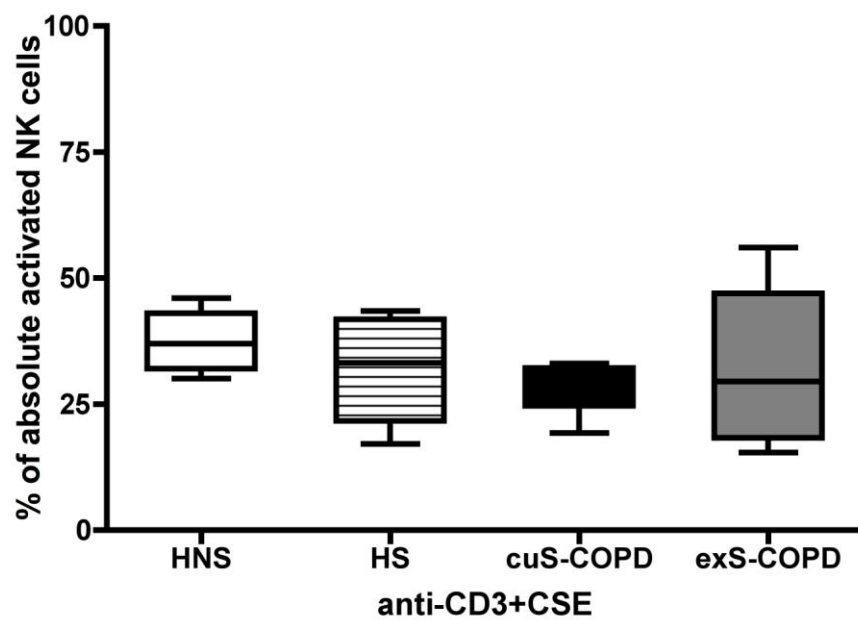


Figure 2. Comparison of absolute activation levels of NKT-like cells upon different stimulation in four groups including healthy non-smokers (HNS, n=4), current healthy smokers (HS, n=4), current smokers with COPD (cuS-COPD, n=4) and ex-smokers with COPD (exS-COPD, n=4). PBMCs were collected and incubated with different reagents for 48 hours.

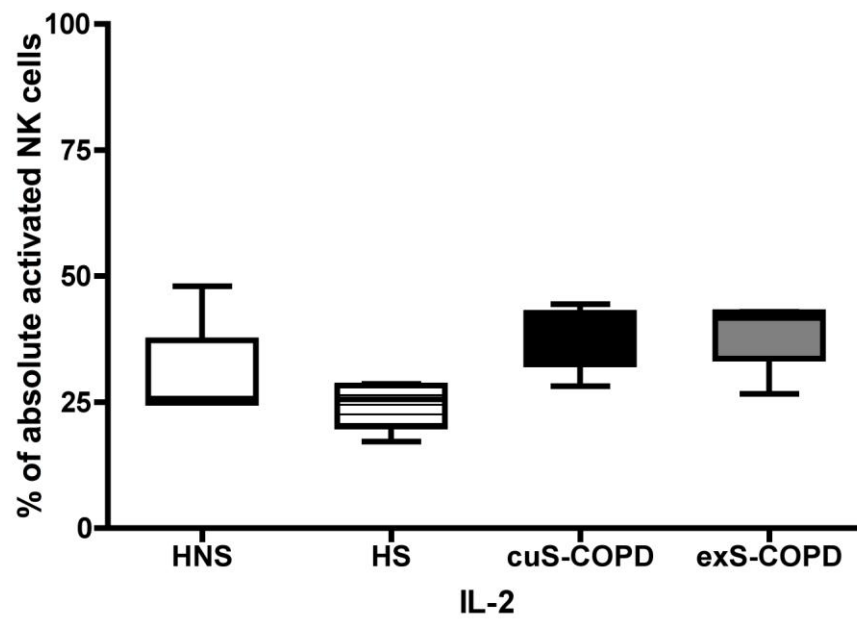
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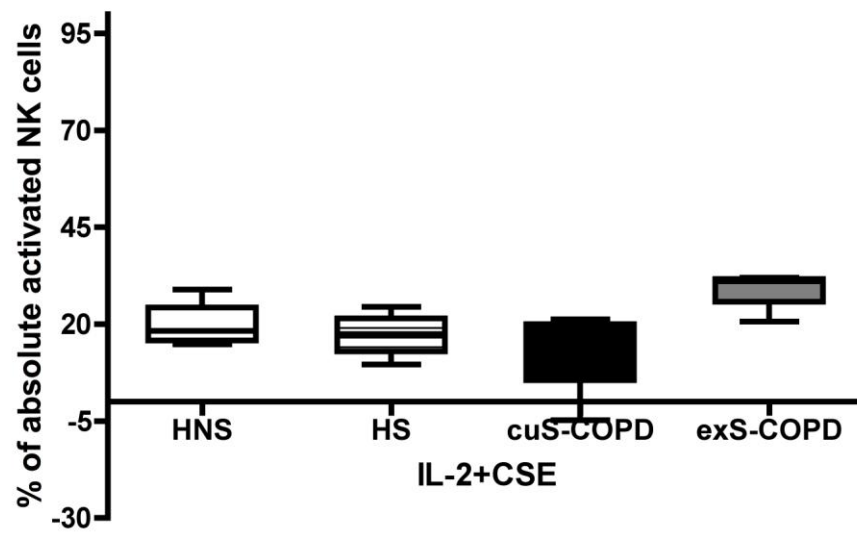
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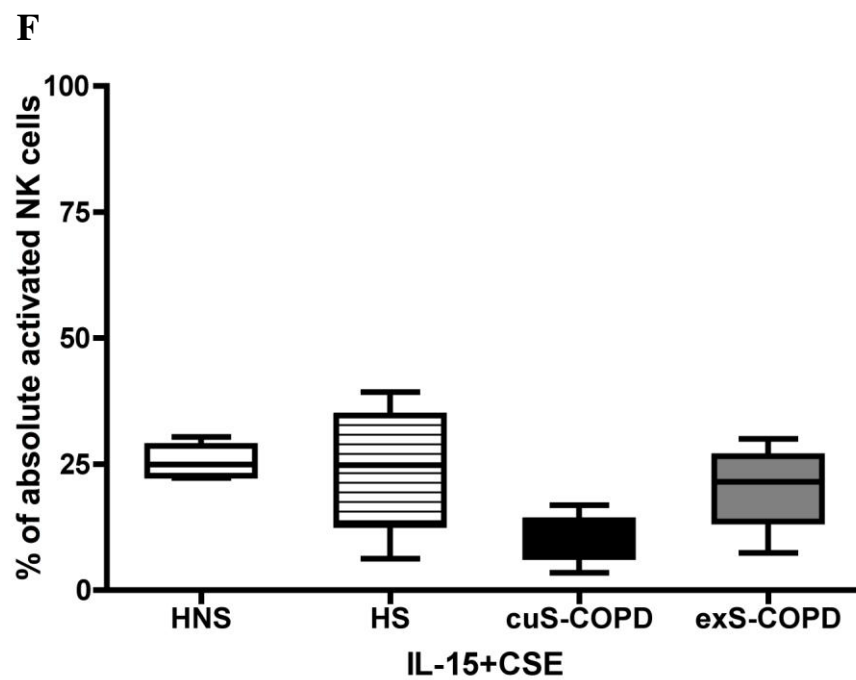
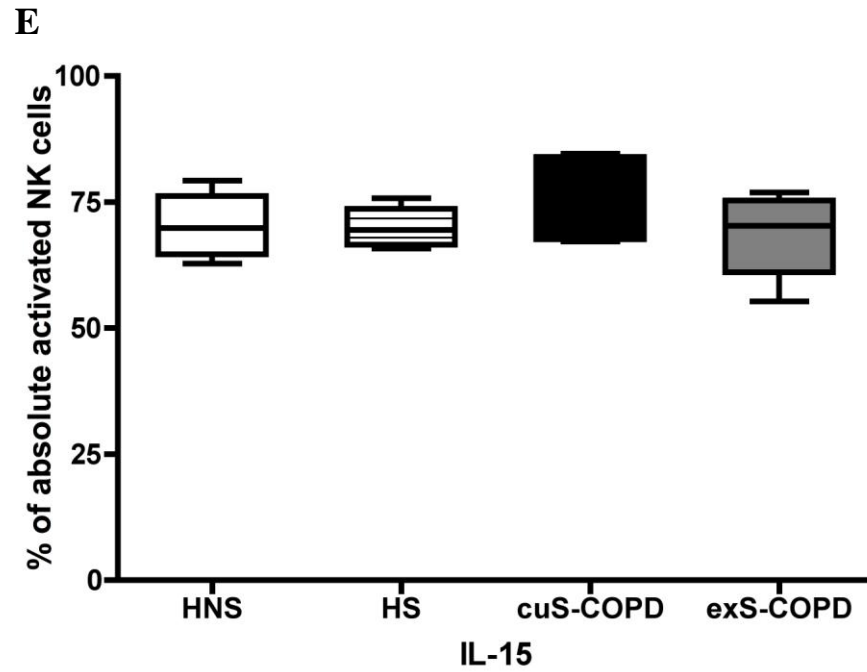


Figure 3. Comparison of absolute activation levels of NK cells upon different stimulation in four groups including healthy non-smokers (HNS, n=4), current healthy smokers (HS, n=4), current smokers with COPD (cuS-COPD, n=4) and ex-smokers with COPD (exS-COPD, n=4). PBMCs were collected and incubated with different reagents for 48 hours.